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Post-processing: Minimum Match 0%
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Perfect score:
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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SUMMARIES

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	AX922612 Sequence	AX355960 Sequence	I12779 Sequence 77	BD197714 Method an	AR311956 Sequence	CQ801520 Sequence	BD254228 Regulatio	BD254227 Regulatio		-	-	AR083535 Sequence	AX316109 Sequence	CS089171 Sequence	AX316096 Sequence	CS089158 Sequence	CQ796912 Sequence	Description	

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ALIGNMENTS

RESULT 2 CS089158 LOCUS DEFINITION ACCESSION VERSION KEYWORDS SOURCE	Query Match Best Local Matches 1 Qy 1 Db 1	VERSION VERSION VERSION VERSION KEYWORDS SOURCE ORGANISM REFERENCE AUTHORS TITLE JOURNAL FEATURES BOURCE ORIGIN	RESULT 1 CQ796912 LOCUS DEFINITION DEFINITION
CS089158 18 bp DNA Sequence 5 from Patent EP1502948. CS089158 CS089158.1 GI:66714442	ch 100.0%; Score 18; DB 6; 1 Similarity 100.0%; Pred. No. 67; 18; Conservative 0; Mismatches 0; 1 CCCCCCTCTTCTGGATAC 18	CQ796912.1 GI:46408538 synthetic construct synthetic construct synthetic construct cother sequences; artificial sequences. Hovanec,T.A. Anmonia-oxidizing bacteria and methods of thesame Patent: WO 2004026772-A 5 01-APR-2004; Aquaria Inc. (US) Location/Qualifiers 118 //organism="synthetic construct" //mol_type="unassigned DNA" //db_xref="taxon:32630" //note="Oligonucleotide probe"	CQ796912 18 bp DNA Sequence 5 from Patent WO2004026772.
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Ammonia-oxidizing bacteria
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Ammonia-oxidizing bacteria
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                                                                                                                                                                                                          Kleesiek,K., Brinkmann,T., Goetting,C. & Xylosyltransferase and isoforms thereof Patent: WO 0149831-A 13 12-JUL-2001; Kleesiek, Knut, Prof. Dr. (DE)
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Sequence 21 from Patent
CQ796928
CQ796928.1 GI:46408554
                                                                                                                  Similarity
                                                                                                                                                                                                                                                                          other sequences; artificial sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Patent:
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llarity 88.2%;
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                                                                                                                                                        /organism="synthetic construct"
/mol_type="unassigned DNA"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   'note="Oligonucleotide probe"
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88.2%;
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17 bp
patent US 6740735.
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Pred. No. 1.5e
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Pred. No. 1.5e+04;
0; Mismatches 2;
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WO0149831.
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B Blatt, L., Zwick, M., Pavco, P. and Mcswiggen, J.

S Blatt, L., Zwick, M., Pavco, P. and Mcswiggen, J.

Regulation of repressor genes using nucleic acid molecules
Patent: JP 2002541795-A 2020 10-DEC-2002;

RIBOZYME PHARMACEUTICALS INC

OS Eukaryote
PN JP 2002541795-A/2020
PD 10-DEC-2002
PP 10-DEC-2002
PP 11-APR-2000 JP 2000611654
PR 12-APR-1999 US 60/129390
PR 12-APR-1999 US 60/129390
PI LAWRENCE BLATT, MICHAEL ZWICK, PAMELA PAVCO, JAMES MCSWIGGEN PC C12N15/09, A61X38/00, A61K48/00, A61P43/00, A61P43/00, C12N5/10, PC
                                                                                                                                                                                                                                                                                                                                                       C12P21/02,
pc
C12P21/02,C12P21/02//A61K31/711,(C12N5/10,C12R1:91),(C12P21/02, PC
C12R1:91),
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Glanturco, S.H. and Bradley, W.A.
DNA encoding human apoB48R: a monocyt
B48 receptor gene and protein
Patent: US 6740735-A 17 25-MAY-2004;
Location/Qualifiers
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Regulation of repressor
BD254227
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                        CCCCCTCTTCTGGA 15
CCCCCTCTCCTGGA 16
                                                               68.9%;
llarity 92.9%;
Conservative
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(C12N5/00,C12R1:91)
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                                                                                                                                                                                                 /organism='Bukaryote'
Location/Qualifiers
                                                                                                                                               /organism="unidentified"
/mol_type="genomic DNA"
/db_xref="taxon:32644"
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/mol_type="genomic
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Pred. No. 9.2e
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BD254228
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OS EUKATYOLE
PN J 2002541795-A/2021
PD 10-DEC-2002
PP 11-APR-2000 JP 2000611654
PR 12-APR-1999 US 60/129390
PR 12-APR-1999 US 60/129390
PI LAWRENCE BLATT, MICHAEL ZWICK, PAMELA PAVCO, JAMES MCSWIGGEN C12N15/09, A61K38/00, A61K48/00, A61P43/00, A61P43/00, C12N5/10, PC C12P21/02,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 (bases 1 to 17)
Blatt, L., Zwick, M., Pavco, P. and Mcswiggen, J.
Regulation of repressor genes using nucleic acid molecules
Patent: JP 2002541795-A 2021 10-DEC-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BD254228.1 GI:330639
JP 2002541795-A/2021.
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Sequence 30 from Patent
CQ801520
CQ801520.1 GI:47058110
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C12P21/02,C12P21/02//A61K31/711,(C12N5/10,C12R1:91),(C12P21/02, PC
C12R1:91),
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                                                                                                                                                            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini;
Hominidae; Homo.
                                                                                Mitchell, J. and de Belleroche, J.
Neurodegnerative disease-associated gene
Patent: WO 2004033723-A 30 22-APR-2004;
IMPERIAL COLLEGE INNOVATIONS LIMITED (GB)
                                                                                                                                                                                                               Homo sapiens
                                                                                                                                                                                                                              Homo sapiens (human)
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(C12N5/00,C12R1:91)
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Location/Qualifiers
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/mol_type="genomic DNA"
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                                                                Location/Qualifiers
/organism="Homo sapiens"
/mol_type="unassigned DNA"
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Pred. No. 9.2e+04;
0; Mismatches 1;
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nucleic acid molecules.
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                                         Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Butheria; Buarchontoglires; Primates; Catarrhini;
Hominidae; Homo.

CE 1 (bases 1 to 17)

Pavco, P.A., Roberts, E., Jarvis, T., Coeshott, C. and Mcswiggen, J.A.
Method and reagent for treating diseases or conditions concerning
molecule participating in vasculogenic response
Patent: JP 2002509721-A 740 02-APR-2002;
RIBOZYME PHARMACEUTICALS INC
OS Homo sapiens (human)
PN JP 2002509721-A/740
PD 02-APR-2002
PF 24-MAR-1999 US 60/079678
PF 24-MAR-1999 US 60/079678
PI JAMES A MCSWIGGEN
PI JAMES A MCSWIGGEN
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A61P29/00.
PC
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AR311956
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2 CCCCCTCTTCTGGA 15
                                                                                                                                                                                                                                                                                                                                                                                BD197714 17 bp RNA linear PAT 17-JUL-200 Method and reagent for treating diseases or conditions concerning molecule participating in vasculogenic response.
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                                                                                                                                                                                                                                                                                                                                        BD197714.1 GI:33007484 JP 2002509721-A/740.
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                              C12N15/09,A61K31/7088,A61K31/7125,A61K48/00,A61P3/10,A61P17/06, PC
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   A61P35/00,A61P43/00,C12N5/10,C12N9/00//A61K35/76,C12N15/00, PC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /organism="unknown"
/mol_type="genomic
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92.9%;
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Pred. No. 9.2e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 12.4; DB 6;
Pred. No. 9.2e+04;
0; Mismatches 1;
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JOURNAL
FEATURES
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AUTHORS
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112779
1.0CUS
DEFINITION
ACCESSION
VERSION
VERSION
KEYWORDS
SOURCE
Search completed: April 7, 2006, 14:36:14 Job time : 940.432 secs
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                                                                                                                         Query Match 67.8%; Score 12.2; DB 6; Length 23; Best Local Similarity 82.4%; Pred. No. 1.2e+05; Matches 14; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                       TITLE
                                                                                                                                                                                                                                                                                                                                                                                                      ORGANISM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 67.8%; Score 12.2; DB 6; Best Local Similarity 82.4%; Pred. No. 1.2e+05; Matches 14; Conservative 0; Mismatches 3;
                                                                                                                                                                                                                                           source
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                                                                                                                                                                                                                                           chain reaction
Patent: US 5427930-A 77 27-JUN-1995;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                      Unknown.

Unclassified.

1 (bases 1 to 23)

1 (kases 1 to 23)

Birkenmeyer, L.G., Carrino, J.J., Dille, B.J., Hu, H.-Y.,

Kratochvil, J.D., Laffler, T.G., Marshall, R.L., Rinehardt, L.A. and
Solomon, N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                I12779 23 bp | Sequence 77 from patent US 5427930. I12779 GI:910161
                                                                                                                                                                                                                                                                                                  Amplification of target nucleic acids using gap filling ligase
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                                                                                   CCCCCTCTTCTGGATAC 18
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C Method and reagent for treating diseases or conditions concerning molecule
C participating in vasculogenic response
H Key
Location/Qualifiers
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Location/Qualifiers
                                                                                                                                                                                                         /organism="unknown"
/mol_type="unassigned DNA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /organism="Homo sapiens"
/mol_type="genomic RNA"
/db_xref="taxon:9606"
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GenCore version 5.1.7
Copyright (c) 1993 - 2006 Biocceleration Ltd.

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Minimum |
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18
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

 66.7 20 11	66.7 20 2 66.7 20 6 66.7 20 10	66.7 19 4 <i>1</i> 66.7 19 10 66.7 20 2 <i>1</i>	7 12 66.7 17 12 8 12 66.7 18 6 A 9 12 66.7 18 12	2 18 100.0 18 12 ADM32708 3 12 66.7 17 8 ABT39927 4 12 66.7 17 8 ACC7980 5 12 66.7 17 10 ADB43132 6 12 66.7 17 10 ADB30251	Result Query No. Score Match Length DB ID 1 18 100.0 18 6 ABA02420
	Aax93192 PCR prime Aas97903 Murine SA Abx13323 Human NSD	Aaf86277 Mutated c Ade10102 DNA encod Aax14630 Trible he	Ado00840 Human mon Aba02421 Type A/A1 Adm32709 Type A AO	Adm32708 Nitrososp Abt39927 Tumour su Acc67980 Murine ol Adb43132 Tumour su Adh30251 Human mon	Description Aba02420 Type A/A1

The invention relates to 4 novel types of ammonia-oxidising bacteria (AOB) found in freshwater aquaria. The bacteria are able to oxidise ammonia to nitrite and are members of the ammonia-oxidising bacteria family of the beta subdivision of Proteobacteria. The 4 types of bacteria can de distinguished on the basis of their 16S rRNA (ribosomal RNA) gene

New bacteria capable of oxidizing ammonia to nitrite, for preventing or alleviating the accumulation of ammonia in fresh water aquaria, seawater aquaria and waste water.

WPI; 2002-075367/10

Hovanec TA, Burrell

S

(AQUA-) AQUARIA INC.

Claim 26; Page 53; 62pp; English.

4.	44	4.	4.	C 41		c 39				3.5	c 3.		ω	ω	3	2.	2	c 2:	2	c 25		c 2		2	c 2
5 11	11	11	2 11	11		9 11										9 11									0 12
61.1	61.1	61.1	61.1	61.1	61.1	61.1	61.1	61.1	61.1	61.1	61.1	61.1	61.1	61.1	61.1	61.1	61.1	61.1	61.1	66.7	66.7	66.7	66.7	66.7	66.7
20	20					18			-		18		17	17	17	17	15	13	13	30	25	25	25	24	20
4 AAS15168	3 AAC73801	2 AAZ02289	2 AAV85631	12 ADM16369	6 AAS98029	12 ADI04227	6 ABK71325	6 ABX00119	6 ABA04522	4 AAF79606	4 AAH78146	3 AAA92441	8 ACC65147	_	8 ACC67173	B ACC63464	3 AAZ44387	5 ABF89342	5 ABF89343	14 ADY03371	10 ADE10105	9 ACH57541	4 AAF86257	6 ABX10559	13 ADR68924
	Aac73801 Mouse IL-	Aaz02289 PCR prime	Aav85631 LRP5 exon	Adml6369 Murine SA	Aas98029 Murine SA	Adi04227 PEAMT1-li		Abx00119 PCR prime	Aba04522 PCR prime	Aaf79606 Human Akt	Aah78146 PCR prime	Aaa92441 MABL-1 H	Murine	Murine		Acc63464 Murine ol			Abf89343 Oligonucl	Ady03371 PCR prime	Adel0105 Mutagenic	Ach57541 DNA targe	Aaf86257 Mutagenic	Abx10559 BRCA1/SMN	Adr68924 Human NSD

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ALIGNMENTS

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RESULT 1
ABA02420
                                                                                                                                                                                                         Type A; type A1; ammonia-oxidising bacterium; AOB; nitrite; 16S rRNA gene; ribosomal RNA; aquarium; aquaculture; waste water treatment; bioremediation; probe; ss.
                                                                                                                                  19-MAY-2000; 2000US-00573684.
                                                                                                                                                17-MAY-2001; 2001WO-US016265.
                                                                                                                                                                                                                                                         29-AUG-2003
04-MAR-2002
                                                                                                                                                               29-NOV-2001.
                                                                                                                                                                              WO200190312-A1.
                                                                                                                                                                                             Nitrosomonadales.
                                                                                                                                                                                                                                         Type A/Al ammonia-oxidising bacterium 16S rRNA gene probe.
                                                                                                                                                                                                                                                                               ABA02420;
                                                                                                                                                                                                                                                                                            ABA02420 standard; DNA; 18 BP
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(first entry)
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ADM31708

ADM31708

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ADM31708

AC ADM3

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Best Local S
Matches 18
                                                            This sequence represents a probe which was used in the detection of 16S rDNA sequence derived from an ammonia oxidising bacteria (AOB). The amplified sequence may be used in a composition which comprises an isolated bacterial strain that oxidizes ammonia to nitrite. The composition may be used for alleviating or preventing the accumulation of ammonia in a medium. The ammonia is reduced by at least 30% when compared with a level of ammonia that would exist in the absence of the bacterial strain. The composition is useful for alleviating or preventing the secural stop of ammonia in aqueous environment, e.g. a freshwater or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           sequences (ABA02416-ABA02419), and are classified as AOB type A (e.g., R7clone140), type A1 (e.g., R7clone187), type B (e.g., R3clone5) and type C (e.g., R3clone47). The invention also encompasses isolated 165 rRNA gene sequences of the ammonia-oxidising bacteria of the invention, oligonucleotide probes and primers for the detection of these bacteria, and compositions comprising the bacteria. The bacteria of the invention are useful in biological filters for reducing ammonia accumulation in both freshwater and seawater aquaria. They may also be used in waste water treatment and in bioremediation processes to reduce the level of pollution caused by ammonia. The present sequence represents a specifically claimed probe for the detection of the 165 rRNA gene sequences of the type A and A1 ammonia-oxidising bacteria (ABA02416-ABA02417). (Updated on 29-AUG-2003 to standardise OS field)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New composition comprising an isolated bacterial strain that oxidizes ammonia to nitrite, useful for alleviating or preventing the accumula
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             19-SEP-2002; 2002US-0386217P.
19-SEP-2002; 2002US-0386218P.
19-SEP-2002; 2002US-0386219P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              of ammonia in aqueous
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                16S rDNA; ammonia oxidising bacteria; AOB; ammonia; nitrite;
aqueous environment; freshwater; seawater; aquarium; ss; probe.
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                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 6;
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                                                                    The invention relates to a novel isolated 17 mer nucleic acid sequence, CC given in the specification, a sequence containing at least 15 consecutive nucleotides from the 17 mer sequence, a sequence with, after optimal that it least 80 % identity to the 17 mer sequence with, after optimal to fany of them under highly stringent conditions, or the complement of any of them, or the corresponding RNA. The novel isolated nucleic cof any of the invention are useful as probes and primers for detecting, identifying, quantifying and/or amplifying a nucleic acid, e.g. as one component of a gene chip, in vitro as (anti)sense resgents, and for production of recombinant polypeptides. Any of the nucleic acids, component of a gene chip, in vitro as (anti)sense resgents and for production of recombinant polypeptides. Any of the nucleic acids, component of a gene chip, in vitro as (anti)sense resgents and for complement of vector or antibodies directed against the polypeptides are useful for preparation of pharmaceuticals for prevention and/or treatment of viral diseases that are characterised by development of tumours or cell degeneration, specifically cancer but also Albeimer's disease and components. Analysis of the expression of the 17 mer nucleic acids in patient samples is useful for disgnosis and/or prognosis of these contains. The polypeptides can also be used to generate antibodies, and both the polypeptide acid sequences of the invention can be used in gene chips. The nucleic acid sequences of the invention can be used in gene
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Cytostatic; virucide; neuroprotective; nootropic; neuroleptic; gene chip; antisense; sense; tumour; cell degeneration; cancer; Alzheimer's disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure; Page 684; 720pp; French
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human fukutin;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Tumour suppression related human fukutin oligo SEQ ID
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens
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                                     This polynucleotide sequence represents a tumour suppression numan fukutin oligonucleotide of the invention
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ne therapy; tumour
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RESULT 5
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Best Local :
                                                                                                                                                                                                                 The present invention relates to murine oligonucleotides (ACC62754-ACC68866), which are associated with tumour suppression, tumour reversion, apoptosis and virus resistance. The Oligonucleotides are useful as (1) as probes and primers for detecting, identifying, quantifying and/or amplifying nucleic acid, e.g. as one component of a gene chip; in vitro as (anti)sense resgents; and (2) for production of recombinant polypeptides. The oligonucleotides are useful for preparation of pharmaceuticals for prevention and/or treatment of viral diseases that are characterised by development of tumours or cell degeneration, specifically cancer but also Alzheimer's disease and schizophrenia
ADB43132;
                                                                                                                                                                                                                                                                                                                                                                                                         New isolated nucleic acid, useful for treating viral diseases associated with tumors and cell degeneration, also related polypeptides, antibodies
                        ADB43132 standard; DNA; 17
                                                                                                                                                                                               Sequence
                                                                                                                                                                                                                                                                                                                                                                    Disclosure; Page 642; 738pp;
                                                                                                                                                                                                                                                                                                                                                                                               and transfected cells.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Telerman A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                17-SEP-2001; 2001FR-00011979.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        tumour suppression; tumour reversion; apoptosis; virus resistance; viral disease; tumour; cell degeneration; cancer; Alzheimer's disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Cytostatic; virucide; neuroprotective; nootropic; neuroleptic; murine; tumour suppression; tumour reversion; apoptosis; virus resistance;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Murine oligonucleotide associated with tumour supression,
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12; Cqnserv
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                                                                                                                                          Conservative
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                                                                                                                                                      66.7%;
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RESULT 6
ADH30251
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AC ADH3
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DT 11-M
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                                                                                                                                                                                                                                                                                                                   The invention relates to the isolation of 6327 nucleotide sequences, and consequence having at least 80% identity, after optimal alignment, with the nucleotides, a sequence that hybridizes under stringent conditions with the nucleotides, or the complement, or corresponding RNA, of the nucleotides. The nucleotides are used as probes or primers for detecting, identifying, quantifying and/or amplifying nucleic acids, as in vitro suppression or reversion, apoptosis and or viral resistance, to produce recombinant polypeptides, and to prepare transgenic animals, as superimental models. The nucleotides (also vectors containing the vectors), the encoded polypeptides and antibodies (cells containing the vectors), the encoded polypeptides and antibodies (Ab) against the polypeptide are useful for prevention and/or treatment or viral infections or diseases characterized by development of tumours or cell degeneration (e.g. Allzheimer's disease or schizophrenia).

Chalysis of the expression of the nucleotides can be used for diagnosis and/or prognosis of these diseases. The nucleotides and polypeptides can be used to screen for their specific interactive molecules, as potentially useful for treating diseases associated with abnormal contentials.
                                                                                                                                                                                                                                          Query Match
Best Local S
               11-MAR-2004
                                               ADH30251;
                                                                                                                                                                                                                                                                                           Sequence
                                                                               ADH30251 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New nucleic acid encoding human prostate membrane-specific antigen, useful e.g. for treatment of tumors and viral infection, also related polypeptide and antibodies.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Disclosure; Page 435; 771pp; French.
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04-DEC-2003
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                                                                                                                                                                                                                                                                                           BP;
                                                                                                                                                                                                                              Conservative
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(first ent
             (first entry)
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                                                                                                                                                                                                                                                                                           5 A; 4 C; 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   nour suppression; tumour reversion; apoptosis;
transgenic animals; Alzheimer's disease; schizophrenia;
                                                                              DNA; 17
                                                                                                                                                                                                                     100.0%; **
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                                                                                                                                                                                                                                           Score 12;
Pred. No.
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                                                                                                                                                                                                                                          DB 10; ]
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RESULT 7
AD000840
ID AD000
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AC AD00
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AC AD00
DT 12-A
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DE Huma
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W Huma
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OS Homx
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Best Local &
                                                                  Human; monocyte-macrophage cell surface apolipoprotein B 48 receptor; apoB4gR; pCR; ss; monocyte; macrophage; reticuloendothelial cell; monocyte adhesion; cardiovascular disease; apoB-mediated cellular uptake; plasma chylomicron; hypertriglyceridaemic triglyceride-rich lipoprotein; ApoE independent pathway; lipoprotein lipase independent pathway; heparin sulfate proteoglycan independent pathway; hypertriglyceridaemia; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        antilipemic; gene therapy; monocyte-macrophage cell surface apoB48 receptor; apoB48R; pattern B phenotype; familial combined hyperlipidaemia; hypercholesterolaemia; hypertriglyceridaemia; hypercholesterolaemia; hypertriglyceridaemia; hupertriglyceridaemia; brimer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention describes a new isolated DNA molecule, encoding a monocyte-macrophage cell surface aposas receptor protein (aposass) comprising a fully defined 1088 amino acid sequence. The DNA molecule is useful in preparing a composition for treating Pattern B phenotype, familial combined hyperlipidaemia, familial or non-familial hypercholesterolaemia, hypertripidyceridaemia or low plasma high-density lipoprotein levels. This sequence represents a primer used in the isolation of human monocyte-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New isolated DNA molecule encoding a monocyte-macrophage cell surface apoB48 receptor protein (apoB48R), useful in preparing a composition treating e.g., familial or non-familial hypercholesterolemia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         06-AUG-1998; 98US-00130242.
31-MAY-2000; 2000US-00583610.
                         Homo sapiens
                                                                                                                                                                                                                                                                                                                                                  12-AUG-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                ADO00840 standard; DNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  macrophage cell surface apoB48 receptor DNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 23; SEQ ID NO 17; 77pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2003-864799/80.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gianturco SH, Bradley
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     12-JUN-2003; 2003US-00459876.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US2003208060-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human monocyte-macrophage cell surface apoB48 receptor primer #5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (GIAN/) GIANTURCO S H. (BRAD/) BRADLEY W A.
                                                                                                                                                                                                                                                                                               monocyte-macrophage cell surface apoB48R cDNA PCR primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      12;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ហ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ហ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CCTCTTCTGGAT 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CCTCTTCTGGAT 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BP; 2 A;
                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             66.7%;
100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 6 C; 4 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 12;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 10; I
. 7.5e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0
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                                                                                                                                                                                                                                                                                                  #6.
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Nitrosomonadales

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ID ABA02421 standard;
                                                                                                                                                                                                                                                                                                                   RESULT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             cc encoding it. The invention also relates to a method of cell-specific conditions it. The invention also relates to a method of cell-specific conditions of therapeutic compounds to human monocytes, macrophages or other reticuloendothelial cells that express the receptor, a method of chihibiting foam cell formation and increased monocyte adhesion to conditions and increased monocyte adhesion to conditions cells and a method of evaluating an individual at risk of a cardiovascular disease. The monocyte-macrophage cell surface apo848 creceptor protein is useful in the apoB-mediated cellular uptake of plasma chylomicrons and remnants and hypertriglyceridaemic triglyceride-rich conditions and remnants and hypertriglyceridaemic triglyceride-rich confident pathway or a heparin sulfate proteoplycan independent confident pathway or a heparin sulfate proteoplycan independent confident to human monocytes, macrophages or other reticuloendothelial confident that express the receptor, for inhibiting foam cell formation and conditional at risk of a cardiovascular disease such as the risk of a cardiovascular disease such as Conditional at risk of a cardiovascular disease such as Conditional conditions and conditional conditions of cDNA encoding the apo848R protein of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Best
                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New isolated monocyte-macrophage cell surface apoB48 receptor protein, useful for cell-specific delivery of therapeutic compounds to human monocytes or macrophages, or for evaluating an individual at risk for cardiovascular disease.
                                                                                               Type A; type A1; ammonia-oxidising bacterium; AOB; nitrite; 16S rRNA gene; ribosomal RNA; aquarium; aquaculture; waste water treatment; bioremediation; PCR primer; 88.
                                                                                                                                                                                                 29-AUG-2003
04-MAR-2002
                                                                                                                                                                Type A/A1 ammonia-oxidising
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 17 BP; 2 A; 6 C; 4 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The invention relates to the human monocyte-macrophage apolipoprotein B 48 receptor (apoB48R) protein and the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gianturco SH,
                                                                                                                                                                                                                                                    ABA02421;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 23; SEQ ID NO 17; 72pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 06-AUG-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local
                                                                                                                                                                                                                                                                                                                                                                                                                                      12;
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                                                                                                                                                                                                                                                                                                                                                                                                   5 CCTCTTCTGGAT 16
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BRADLEY W
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Similarity
                                                                                                                                                                                                                                                                                                                                                                     CCTCTTCTGGAT 16
                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                 (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Bradley
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 98US-00130242
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  D Q
                                                                                                                                                                                                                                                                                     DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                       100.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       66.7%;
                                                                                                                                                                                                                                                                                     18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      .04;
                                                                                                                                                                                                                                                                                     ₽P
                                                                                                                                                                                                                                                                                                                                                                                                                                      0
                                                                                                                                                                  bacterium
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 12;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                  16S rRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 12; 1
7.5e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                      0
                                                                                                                                                                  gene forward
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           cell surface
nucleic acid
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ADM32709/c
ID ADM327
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AC ADM327
AC ADM327
DT 17-JUN
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TYPE A
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H 16S rD
KW 16S rD
KW 20004
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W 20004
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PN 10-SEI
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CC (ADB) found in freshwater aquaria. The bacteria are able to oxidise cammonia to nitrite and are members of the ammonia-oxidising bacteria are able to oxidise cammonia to nitrite and are members of the ammonia-oxidising bacteria are able to oxidise cammonia to nitrite and are members of the ammonia-oxidising bacteria are able to oxidise cammonia to nitrite and are members of the interval are also and types of bacteria are distinguished on the basis of their 165 rRNA (ribosomal RNA) gene are gequences (ABA02416-ABA02419), and are classified as AOB type A (e.g., R7clone140), type A1 (e.g., R7clone187), type B (e.g., R3clone5) and type CC (e.g., R3clone47). The invention also encompasses isolated 165 rRNA (ribosomal representation) are useful in biological filters for the detection of these bacteria, are described and compositions comprising the bacteria. The bacteria of the invention are useful in biological filters for reducing ammonia accumulation in both freshwater and seawater aquaria. They may also be used in waste water treatment and in bioremediation processes to reduce the level of pollution caused by ammonia. Sequences ABA02421-ABA02422 represent PCR conditions for the detection of the 165 rRNA gene sequences of the type A and A1 ammonia-oxidising bacteria (ABA02416-ABA02417). (Updated on 29-AUG cond A1 ammonia-oxidising bacteria (ABA02416-ABA02417).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local S
Matches 12
                                                                   19-SEP-2002; :
19-SEP-2002; :
19-SEP-2002; :
                                                                                                                                                                                                                                                                                                                                                                                                                                  16S rDNA; ammonia oxidising bacteria; AOB; ammonia; nitrite;
aqueous environment; freshwater; seawater; aquarium; ss; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New bacteria capable of oxidizing ammonia to nitrite, for preventing alleviating the accumulation of ammonia in fresh water aquaria, seawa
         (AQUA-) AQUARIA INC
                                                                                                                                                                                         10-SEP-2003; 2003WO-US028210
                                                                                                                                                                                                                                                      01-APR-2004.
                                                                                                                                                                                                                                                                                                                                                                            Nitrosomonas aestuarii.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Type A AOB 16S rDNA forward primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  17-JUN-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADM32709 standard; DNA; 18 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2002-075367/10.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     l Similarity
12; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        standardise OS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Page 10; 62pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TCTTCTGGATAC 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           wäste water.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                               2002US-0386217P.
2002US-0386218P.
2002US-0386219P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Burrell
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        7 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        5 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 12; DB 6; I
Pred. No. 7.5e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity
Matches 12; Conserv
                                                                     New nucleic acid, useful for p treatment of genetic diseases,
                                                                                                                                                                                                                                                                                                                               Triplex; cis-polypurine tract; cPPT; central termination sequence; CTS; cis acting region; retrovirus; blood disorder; developmental disorder; brain disorder; nervous system disorder; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      composition may be used for alleviating or preventing the accumulation of ammonia in a medium. The ammonia is reduced by at least 30% when compared with a level of ammonia that would exist in the absence of the bacterial strain. The composition is useful for alleviating or preventing the accumulation of ammonia in aqueous environment, e.g. a freshwater or
                                   Example
                                                                                                          WPI; 2001-273782/28.
                                                                                                                                                                                             12-OCT-1999;
                                                                                                                                                                                                                      10-OCT-2000; 2000WO-EP010418
                                                                                                                                                                                                                                              19-APR-2001.
                                                                                                                                                                                                                                                                                              Synthetic.
                                                                                                                                                                                                                                                                                                         Human immunodeficiency virus 1.
                                                                                                                                                                                                                                                                                                                                                                                  Mutated
                                                                                                                                 Charneau P,
                                                                                                                                                         (INSP ) INST PASTEUR.
                                                                                                                                                                                                                                                                     WO200127304-A2
                                                                                                                                                                                                                                                                                                                                                                                                           24-JUL-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                   AAF86277
                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAF86277 standard; DNA; 19 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              seawater aquarium.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   This sequence represents a primer which was used in the detection of the sequence derived from an ammonia oxidising bacteria (AOB). The amplified sequence may be used in a composition which comprises an isolated bacterial strain that oxidizes ammonia to nitrite. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 7; Page 34; 98pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        of ammonia in aqueous environment.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New composition comprising ammonia to nitrite, useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2004-304936/28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Hovanec
                                                           central termination
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TCTTCTGGATAC 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      BP; 7 A; 4 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                Zennou V,
                                                                                                                                                                                                                                                                                                                                                                                  fragment cPPT-225
                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                               99US-0158387P
                              81pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            66.7%;
                                                         sequence
                                                                                                                                 Pflumio F,
                                                                     for prophylactic, ameliorative eases, comprises retroviral cis-
                                 English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      5 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     an isolated bacterial strain that for alleviating or preventing the
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                                                                                                                                 Dubart Kupperschmitt
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This invention relates to a triple stranded nucleic acid containing at least one copy of the cis-polypurine tract

(cppT)

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RESULT 11
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        central termination sequence (CTS) cis acting regions of a retrovirus. The CPPT region initiates synthesis of a downstream plus strand during reverse transcription and CTS is involved in the termination and ejection of reverse transcriptase. The CPPT and CTS regions induce a 3-stranded
                                                                                                                                            (ZENN/)
(PFLU/)
(SIRV/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence
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                                                                                                                                                                                                                                                                                                                                                                                     10-APR-2002; 2002US-00122114.
                                                                                                                                                                                                                                                                                                                                                                                                                                                             16-OCT-2003.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human immunodeficiency virus
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                                                                                                        ) CHARNBAU P.
) ZENNOUŤ V.
) PELUMIO P.
) PELUMIO P.
) SIRVEN A.
) DUBART KUPPERSCHMITT A
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /note= "This sequence lacks both start and stop codons"
/transl_except= (pos:1,aa:Asn)
/note= "This codon has an apparent 2 nucleotide deletion
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Location/Qualifiers
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Pred. No
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Zennou V,

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Sirven A,

Dubart Kupperschmitt

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The present sequence represents a potential triple-helix forming region. It can be used to demonstrate the assay of the invention. The assay comprises adding a sample containing double-stranded DNA test sequences, e.g. containing the present sequence, to an aqueous medium containing at least one complex of anchor DNA, attached to a solid support, and reporter DNA, where either a part of the anchor DNA or reporter DNA is

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SXCCCCCCCCX SX PPT PTX RR
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present invention relates to the isolation of a lentiviral polymucleotide (e.g. HIV-1) polymurine tract cis-active (CPPT) and central termination sequence (CTS) cis-acting regions that induce a three stranded or triplex DNA structure. The lentiviral polymucleotide sequences are useful in gene therapy for preparing a composition for treating an individual suffering from, or having a likelihood of developing a disorder such as brain or developmental disease. The present sequence encodes a mutant HIV-1 CPPT peptide.
                                                                                                                                           Assay of genetic sequences based on triplex formation from double stranded analyte - and hybrid of anchor and reporter sequences, w reporter released if triplex formation occurs, used e.g. to ident
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New nucleic acid comprising at least one copy of the cPPT and CTS cisacting regions of a retrovirus that induce a three-stranded DNA structure, useful for preparing a composition for treating e.g., brain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2003-844442/78.
P-PSDB; ADE10101.
                                                                                                                                                                                                                                                                                                        22-DEC-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                Triple-helix fo 
identification;
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Triple helix forming nucleotides 2407-2426 of the n-myc gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   24-MAR-1999
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                                                                                                      Disclosure; Col 13-14; 168pp; English
                                                                                                                                                                                                  WPI; 1999-130384/11.
                                                                                                                                                                                                                                                                                29-OCT-1992;
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                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                forming region; Triplex formation; on: bacteria; oncogene; virus; ds.
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Pred. No.
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7.5e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                              DNA detection;
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RESULT 13
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                                                   Query Match
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                                        Matches
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Best Local
                                                                                                            AAX91991-X97517 represent PCR primers used to amplify open reading frames and other nucleic acid sequences from the genome of Chlamydia pneumoniae (see AAX91990). C. pneumoniae causes respiratory disease such as pneumonia and bronchitis and is thought to be a contributing factor in heart disease, sarcoidosis, sinusitis, purulent otitis media, erythema nodosum or pharyngitis. The polypeptides encoded by the open reading frames of the C. pneumoniae genome (see AAX94584- AAX95879) can be used in immunogenic compositions as vaccines. Vectors containing C. pneumoniae esquences can also be used as immunogenic compositions, especially where the vector directs the expression of a neutralising epitope of C. pneumoniae
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Respiratory disease; pneumonia; bronchitis; heart disease; sarcoidosis; sinusitis; purulent otitis media; erythema nodosum; pharyngitis; vaccin neutralising epitope; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  which is detected as an indication of the presence of the DNA test which is detected as an indication of the presence of the DNA test sequence. The method is used to detect DNA sequences, particularly for identification of bacteria (by detecting genes for ribosomal RNA) in clinical samples, but also detection of oncogenes and Hepatitis B virus
                                                                                       Sequence
                                                                                                                                                                                                                                                                                              Genome
                                                                                                                                                                                                                                                                                                                                                                                              21-NOV-1997;
04-NOV-1998;
                                                                                                                                                                                                                                                                  Page 1570; Disclosure; 1912pp;
                                                                                                                                                                                                                                                                                                                     WPI; 1999-357842/30
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                                                                                                                                                                                                                                                                                                                                                                  (GEST ) GENSET:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Chlamydophila
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PCR primer used
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               s
                                                                                                                                                                                                                                                                                          sequence of Chlamydia pneumoniae.
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                                      . Similarity 12; Conserv
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CCTCTTCTGGAT 16
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98US-0107078F
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RESULT 14
AAS97903/c
ID AAS97903 standard; DNA;
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                                                                                                                                                                                                                                          Bachmanov AA, be
                                                                                                                                                                                         Novel isolated polypeptide comprising variant form of mouse or human SAC polypeptide, and is associated with altered preference for carbohydrates or other sweeteners, useful for preventing obesity, diabetes, alcoholism
                                                                                                                                                                                                                                                                                                                                                                                                               blood; to
                                                                                                                                                                         Claim 14; Page 92; 239pp; English.
                                                                                                                                                                                                                                                                                               28-APR-2000; 2000US-0200794P
28-JUL-2000; 2000US-0221419P
10-NOV-2000; 2000US-0247443P
                                                                                                                                                                                                                              WPI; 2002-075162/10.
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                                                                                                                                                                                                                                                                                                                                                                                     Mus sp.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAS97903;
                                                                                                                                                                                                                                                                                                                                                                                                             mouse; SAC1; carbohydrate; sweetener; ethanol; alcoholism; ss;
y; diabetes; transgenic embryo; body tissue; body fluid; pancreas;
tongue; PCR primer; anorectic; antidiabetic; gene therapy;
                                                                                                                                                                                                                                                                       WARNER LAMBERT CO.
MONELL CHEM SENSES CENT.
                                                                                                                                                                                                                                                                                                                                                                                                      replacement
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1 DR, Ross D,
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The invention relates to an isolated polypeptide, comprising a variant CC form of mouse or human SAC1 polypeptide. The variant form is associated CC with altered preference for carbohydrates, other sweeteners or ethanol. CC The polypeptide and its associated DNA sequence can be produced by CC recombinant techniques and is useful for preventing obesity, diabetes or CC alcoholism associated with SAC1 expression. The sequences are useful in CC screening for drugs and sweeteners. Recombinant cell lines and transgenic CC embryos may be used in screening for and identifying agents that induce CC or repress function of SAC1. Predisposition to diabetes, obesity or CC cloholism can be ascertained by testing any fluid or tissue of a human CC (such as blood, pancreas or tongue) for sequence variations of the SAC1 CC gene. A sequence variation of the SAC1 locus may indicate a CC diagnostic mark. The polynucleotide can be detected in a biological CC sample by contecting the DNA with a probe to form a hybridisation complex which is then detected. The sequences represent cDNA encoding human and course SAC1 polypeptides and PCR primers specific for the SCA1 genes

Sequence 20 B₽; 7 ? σ Ç σ ဓ μ. H '' 0 u; 0 Other;

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                                  Similarity
CCCTCTTCTGGA 8
             CCCTCTTCTGGA 15
                            Conservative
                         100.08;
                                         66.7%;
                                  Score 12;
Pred. No.
                            Mismatches
                                  DB 6; Lo. 7.5e+03;
                                        Length 20;
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Search completed: April 7, 2006, 14:43:39 Job time: 216.568 secs
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ABX13323/c
                                                                                                                                                                                                                                                                             The invention relates to diagnosing congenital hemidysplasia, CC ichthyosiform erythroderma and limb defects (CHILD) syndrome or psoriasis CC in a patient, involves detecting nucleotide difference between patient CC NSDHL oligonucleotide (gene encoding 3beta-hydroxysteroid dehydrogenase CC (3beta-HSD)) and wild type NSDHL gene, or detecting accumulation of CC sterol intermediates before the generation of cholest-8(9)-en-3beta-01 in CC the cholesterol biosynthetic pathway, in the body fluids or cells of the CC patient. The method is useful for diagnosing CHILD syndrome and psoriasis CC in a patient. Other diseases implicated in defects of the cholesterol CC biosynthetic pathway include mevalonic aciduria, desmosterolosis, Smith-CC temil-Opitz syndrome (SLOS), Conradi-Hunermann-Happle syndrome; CC chondroplasia punctata (X-linked disease), bone disorders, osteoporosis, CC condeceptosis, skin disorders, ichthyosis, eye disorders, (e.g. cataract CC and microopthagmia) and arthritis. The human NSDHL gene is located on CC chromosome X128. The present sequence is a primer for routine sequencing CC of and mutation detection in the human NSDHL gene (one of 8 exons or the
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Best Local S
Matches 12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Diagnosing CHILD syndrome or psoriasis, by detecting differences between patient and wild type genes encoding 3 beta-hydroxysteroid dehydrogenase or accumulation of sterol intermediates in body fluids of the patient.
                                                                                                                                                                                                                              Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        01-JUN-1999; 99US-0137020P.
01-JUN-2000; 2000US-00588976.
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skin disorder; primer.
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1: /cgn2_6/ptodata/1/ina/1_COMB.seq:*

2: /cgn2_6/ptodata/1/ina/5_COMB.seq:*

3:\$\'cgn2_6/ptodata/1/ina/6A_COMB.seq:*

4: /cgn2_6/ptodata/1/ina/6B_COMB.seq:*

5: /cgn2_6/ptodata/1/ina/PCTUS_COMB.seq:*

6: /cgn2_6/ptodata/1/ina/PCTUS_COMB.seq:*

7: /cgn2_6/ptodata/1/ina/PCOMB.seq:*

8: /cgn2_6/ptodata/1/ina/RE_COMB.seq:*

9: /cgn2_6/ptodata/1/ina/RE_COMB.seq:*
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Copyright (c) 1993 - 2006 Biocceleration Ltd
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US-09-198-452A-2493
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Sequence 17, Appl Sequence 2493, App Sequence 24935, A Sequence 4, Appli Sequence 14, Appli Sequence 14, Appli Sequence 154, App Sequence 126, App Sequence 126, App Sequence 126, App Sequence 126, App Sequence 24, Appli Sequence 24, Appli Sequence 23, Appli Sequence 23, Appli Sequence 10385, Appli Sequence 10386, Appli Sequence 10386, Appli Sequence 10387, Appli Sequence 
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ALIGNMENTS	US-08-929-856-58	US-08-311-486C-127	US-08-311-486C-126	US-08-311-486C-125		96-196G-	US-09-396-196G-103141	US-09-396-196G-103140	US-09-396-196G-100180	US-09-396-196G-100179	US-09-396-196G-86459	US-09-396-196G-84565	US-09-396-196G-84554	US-09-396-196G-84553	US-09-396-196G-66069	US-09-396-196G-62035	US-09-396-196G-52911	US-09-396-196G-52017	US-09-396-196G-52016	US-09-396-196G-52015	US-09-396-196G-52014	
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	58, Appl	127, App	126, App	125, App	4, Appli	117806,	103141,	103140,	100180,	100179,			84554, A			62035, A	•	•	•	•	52014, A	

US-09-583-610D-17 APPLICANT: Gianturco, S.H. APPLICANT: Bradley, W.A. TITLE OF INVENTION: DNA Encoding Human apoB48 TITLE OF INVENTION: Apolipoprotein B48 Recep FILE REFERENCE: D5880CIP CURRENT APPLICATION NUMBER: US/09/583,610D CURRENT FILING DATE: 2000-05-31 PRIOR APPLICATION NUMBER: US 09/130,242 PRIOR FILING DATE: 1998-08-06 RESULT 1 US-09-583-610D-17 Query Match Best Local S Matches 12 Patent No. 6740735 GENERAL INFORMATION: NUMBER OF SEQ ID NOS: 22 SEQ ID NO 17 LENGTH: 17 TYPE: DNA Sequence 17, Application US/09583610D Patent No. 6740735 OTHER INFORMATION: Antisense primers were based on the 5'-end OTHER INFORMATION: sequence of the THP-1 (73 clone (bp 2203-2187 OTHER INFORMATION: of ApoB48R cDNA) ORGANISM: Artificial Sequence FEATURE: 12; 5 CCTCTTCTGGAT 16 5 CCTCTTCTGGAT 16 Similarity Conservative 100.0%; --Score 12; DB ; Pred. No. 1e+ 0; Mismatches apoB48R: A Monocyte-Macrophage Receptor Gene and Protein DB 3; 1e+03; 0 Length 17; Indels 0 Gaps

RESULT 2 US-08-173-489C-17/c

Sequence 17, Application US/08173489C
Patent No. 5861244
GENERAL INFORMATION:
APPLICANT: WANG, C. -G.
APPLICANT: WANG, C. -G.
APPLICANT: HEPBURN, A. G.
TITLE OF INVENTION: TRIPLE-STRAND FORWATION.
NUMBER OF SEQUENCES: 365

CORRESPONDENCE ADDRESS:
ADDRESSEE: PROFILE DIAGNOSTIC SCIENCES, INC.,
STREET: 510 EAST 73RD STREET,

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ZIP: 10021 COUNTRY:

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Sequence 2493, Application US/09198452A

Patent No. 6559294

GENERAL INFORMATION:
APPLICANT: Griffais, R.

TITLE OF INVENTION: Chlamydia pneumoniae genomic sequence and polypeptides, fragments
TITLE OF INVENTION: thereof and uses thereof, in particular for the diagnosis, preve
TITLE OF INVENTION: and treatment of infection
TITLE OF INVENTION: and treatment of infection
FILE REFERENCE: 9710-003-999
CURRENT APPLICATION NUMBER: US/09/198,452A
CURRENT FILING DATE: 1998-11-24
NUMBER OF SEQ ID NOS: 6849

SEQ ID NO 2493
LENGTH: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ORIGINAL SOURCE:
ORGANISM: Homo sapiens
PUBLICATION INFORMATION:
AUTHORS: Ibson, J M, Rabbitts, P H.
TITLE: Sequence of a germ-line N-myc
Patent No. 5861244
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local (
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APPLICATION NUMBER: US 07/968,436

PILING DATE: 29 OCT 1992

ATTORNEY/AGENT INFORMATION:

NAME: Handelman, Joseph H.

REGISTRATION NUMBER: 26,179

REGISTRATION NUMBER: U9518-6

TELECOMMUNICATION INFORMATION:

TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEPHONE: (attorney) (212) 708-1880
TELEPAX: (attorney) (212) 246-8959
INFORMATION FOR SEQ ID NO: 17:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM PC/XT/AT
COMPUTER: IBM PC/XT/AT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: Nucleic Acid
STRANDEDNESS: double
TOPOLOGY: linear
MOLECULE TYPE: Genomic DNA
DESCRIPTION: n-myc gene (Accession # Y00664)
DESCRIPTION: nucleotides 2407 to 2426
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER: IBM PC/XT/AT
OPERATING SYSTEM: MS-DOS version 6.2
SOFTWARE: Wordperfect Version 5.1
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   HYPOTHETICAL:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ANTI-SENSE: NO
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Local Similarity 100.0%;
108 12; Conservative (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PAGES: 399-
DATE: 1988
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  VOLUME:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TITLE: gene and amplification as a mechanism of TITLE: activation
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    JOURNAL:
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b; Pred. No. 1.1
0; Mismatches
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thes 0;
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4 CCCTCTTCTGGA 15

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CURRENT APPLICATION NUMBER: US/09/396,196G
; CURRENT FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: 60/100,678
; PRIOR FILING DATE: 1998-09-17
; NUMBER OF SEQ ID NOS: 127806
; SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 24325
; ELNCTH: 25
; TYPE: DNA
; ORGANISM: Mus musculus
US-09-396-196G-24325
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Best Local Similarity 1
"""ches 12; Conservati
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; TYPE: DNA
; ORGANISM: Chlamydia pneumoniae
US-09-198-452A-2493
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-10-002-623-451; Sequence 451, Application Patent No. 6929911; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       밁
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; ORGANISM: Homo Sapiens
US-10-002-623-451
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                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:

APPLICANT: Michael Mittmann
APPLICANT: David Mack
APPLICANT: David Lockhart
APPLICANT: David Lockhart
APPLICANT: Affymetrix, Inc.
TITLE OF INVENTION: Methods of Genetic Analysis
FILE REFERENCE: 3101.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQ ID NOS: 952
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 451
LENGTH: 20
Query Match
Best Local Similarity
Matches 12; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 24325, Application US/09396196G Patent No. 6821724
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CURRENT APPLICATION NUMBER: US/10/002,623
CURRENT FILING DATE: 2001-11-01
PRIOR APPLICATION NUMBER: US 60/245,355
PRIOR FILING DATE: 2000-11-01
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: OEFNER, PETER J.
APPLICANT: UNDERHILL, PETER A.
TITLE OF INVENTION: A METILIATION, SUBSTRUCTURE AND GENE FLOW WITHIN HUMAN
TITLE OF INVENTION: APPLILATION, SUBSTRUCTURE AND GENE FLOW WITHIN HUMAN
TITLE OF INVENTION: POPULATIONS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: DNA
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12; Conserv
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66.7%; Score 12; DB 3; Liarity 100.0%; Pred. No. 1.1e+03; Conservative 0; Mismatches 0;
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                                      Length 25;
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  Gaps
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RESULT 7
US-09-474-922A-14
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               GENERAL INFORMATION:
APPLICANT: Bre
APPLICANT: Lex M.
                                                               Sequence 14, Application US/09474922A Patent No. 6187586
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 4, Application US/09094714A Patent No. 6117847
                                                                                                                                                                                                                                 Matches
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                                                                                                                                                                                                                                                               Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
APPLICANT: C. Frank Bennett, Nicholas M. Dean
TITLE OF INVENTION: OLICONUCLECTIDES FOR ENHANCED MODULATION
TITLE OF INVENTION: PROTEIN KINASE C EXPRESSION
                                                                                                                                                                                                                                                                                                                                                                                               TELEFAX: (215) 568-3439 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE, DISKETTE,
COMPUTER: IBM PS/2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/089,996
FILING DATE: 09-JUL-1993
PRIOR APPLICATION DATA:
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APPLICATION NUMBER: US/09/094,714A
FILING DATE: June 15, 1998
                                                                                                                                                                                                                                                                                                                                                                                                                                           REFERENCE/DOCKET NUMBER: IS
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: 07/852,852 FILING DATE: 16-MAR-1992 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS:
ADDRESSE: Woodcock Washburn Kurtz Mackiewicz & No. 6117847ris, LLP STREET: One Liberty Place - 46th Floor
                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE: 07-JUN-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/601,269
PILING DATE: 14-FEB-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                              STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME: Paul K. Legaard
REGISTRATION NUMBER: 38,534
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: WORDPERFECT 8.0
                                                                                                                                                                                                                                                                                                                                                                                                                              TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              OPERATING SYSTEM: PC-DOS
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                                                                                                                                                                                                                                                                                                                                            nucleic acid
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          Brett P. Monia
Lex M. Cowsert
                                                                                                                                                                CCCCCTCTTCT 12
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Richard A. Roth
                                                                                                                                                                                                                                 Conservative
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                                                                                                                                                                                                                                              Score 11;
; Pred. No.
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                                                                                                                                                                                                                                                              DB 3;
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                                                                                                                                                                                                                                                              Length 15;
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US-09-280-799-154
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                                                                                                                                                                                                                                                                                                                                  US-09-277-020-18
                                                                                                                                                                                                                                                                                                                                                    RESULT 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILE REPERENCE: ISH-0340
CURRENT APPLICATION NUMBER: US/09/280,799
CURRENT FILING DATE: 1999-03-26
NUMBER OF SEQ ID NOS: 208
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 154
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
APPLICANT: Dean, Nicholas M.
APPLICANT: Karras, James G
APPLICANT: Kokay, Robert
TITLE OF INVENTION: ANTISENSE MODULATION OF INTERLEUKIN-5 SIGNAL
TITLE OF INVENTION: TRANSDUCTION
                                    SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 18
LENGTH: 20
                                                                                                                                                                                                                                                     GENERAL INFORMATION:
APPLICANT: Bennett, C. Frank
                                                                                                                                                                                                                                                                                           Sequence 18, Application US/09277020 Patent No. 6210892
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQ ID NOS:
SEQ ID NO 14
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Best Local :
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                                                                                                              TITLE OF INVENTION: Alteration of Cellular Behavior by Antisense Modulation TITLE OF INVENTION: of mRNA Processing FILE REFERENCE: ISPH-0339 CURRENT APPLICATION NUMBER: US/09/277,020 CURRENT FILING DATE: 1999-03-26 EARLIER APPLICATION NUMBER: 09/167,921 EARLIER FILING DATE: 1998-10-07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION NUMBER: US/09/474,922A
CURRENT FILING DATE: 1999-12-29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: FILE REFERENCE:
                                                                                              NUMBER OF SEQ ID NOS: 65
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 20
TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LENGTH: 18
TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     OTHER INFORMATION: Description of Artificial Sequence: Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ORGANISM: Artificial Sequence FEATURE:
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Similarity 100.0%; Pred. No. 3.7e+03;
11; Conservative 0; Mismatches 0;
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; Pred. No. 3.8e+03;
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US-09-060-299-126
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; OTHER INFORMATION: Description of Artificial Sequence:Synthetic
US-09-277-020-18
                                                                                                                              US-09-060-299-126
                                                                                                                                                               IBLEFAX: (703)816-4100
INFORMATION FOR SEQ ID NO: 126:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: nucleic acid
                                                                                 Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 126, Application US/09060299 Patent No. 6545137
                                                                    Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
                                                                                                                                                                                                                                                                                                                        CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 60/043,553
FILING DATE: 15-APR-1997
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 60/048,740
FILING DATE: 05-JUN-1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Patent No. 6545137
GENERAL INFORMATION:
APPLICANT: Todd,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                NAME: B.J.Sadoff
REGISTION NUMBER: 36,663
REFERENCE/DOCKET NUMBER: 620
TELECOMMUNICATION INFORMATION:
TELEPHONE: (703)816-4091
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: PATENTIN Release #1.0, Version #1.25 (EPO)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/060,299
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CITY: Arlington
STATE: Virginia
                                                                                                                                                         STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE: 15-APR-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Local
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                                     IJ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           T: Kawaguchi, Yoshihiko
T: Merriman, Tony R
T: Metzker, Michael L
INVENTION: No. 6545137el Receptor
                                CCTCTTCTGGA 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CCCTCTTCTGG 20
         CCTCTTCTGGA 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Virginia
: US
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1100 No. 6545137th Glebe Road, Eighth Floor
                                                            Conservative 0
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Hammond, Holly
Hey, Patricia
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Hess, John W
Caskey, Charles T
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               455
                                                                                                                                                                                                                                                                                   620-35
                                                                                 Score 11; DB 3; Pred. No. 3.8e+(
                                                                   0;
                                                                    Mismatches
                                                                   . 3.8e+03;
ches 0;
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                                                                   Indels
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                                                                   Gaps
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RESULT 12
US-08-117-952-432/c
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US-09-402-923A-126
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GENERAL INFORMATION:
                                                                                                                                                                                                         Query Match
                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                                                                                                                              INFORMATION FOR SEQ ID NO: 126: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PILING DATE: 14-Feb-2001

PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/GB98/01102
PILING DATE: 15-APR-1998
APPLICATION NUMBER: US 60/043,553
FILING DATE: 15-APR-1997
APPLICATION NUMBER: US 60/048,740
PILING DATE: 05-UN-1997
                                                                                                                                                                                        Local
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APPLICATION NUMBER: US/09/402,923A
                                                                                                                                                                                                                                                                                                                                                                                                                                                          REGISTRATION NUMBER: 36,663
REFERENCE/DOCKET NUMBER: 620-81
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COUNTRY: US
ZIP: VA 22201-4714
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Kawaguchi, Yoshihiko
Merriman, Tony R
Metzker, Michael L
TITLE OF INVENTION: NO. 6555654el LDL-Receptor
NUMBER OF SEQUENCES: 455
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ATTORNEY/AGENT INFORMATION: NAME: B.J.Sadoff
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Todd, John A
Hess, John W
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                                                                                                                                                                    h 61.1%; Score 11; DB 3; Louis 11; Score 11; DB 3; Louis 11; Similarity 100.0%; Pred. No. 3.8e+03; 11; Conservative 0; Mismatches 0;
                                                                                    CCTCTTCTGGA 18
                                                                                                                                                                                                                                                                                                           LENGTH: 20 base pairs
TYPB: nucleic acid
STRANDEDNESS: single
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                                                                                                                                                                                                                                                                                                                                                                                                                     TELEPHONE: (703)816-4091
TELEPAX: (703)816-4100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CITY: Arlington
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Hammond, Holly
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                                                                                                                                                                                                            Length 20;
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Sequence 432, Application US/08117952 Patent No. 5851760 GENERAL INFORMATION: APPLICANT: Evans, Glen A. APPLICANT: Smith, Michael W.

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NUMBER OF SEQ ID NOS: 35
SOPTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 18
LENGTH: 22
TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                Sequence 18, Application US/10341199 Patent No. 6852494
                                                                                                                                                                                                                                                                                                                                                               GENERAL
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                                                                                                                                                                                                                                                                                                                                                                                                                       10-341-199-18/c
                                                                                                                  FILE REFERENCE: 12251-036001
CURRENT APPLICATION NUMBER: US/10/341,199
CURRENT FILING DATE: 2003-01-10
                                                                                                                                                                     APPLICANT: Huang, Yih
APPLICANT: Shen, Ming
APPLICANT: Law, Simon W.
APPLICANT: Huang, Tai-Nang
TITLE OF INVENTION: NUCLBIC ACID AMPLIFICATION
                                                                                                                                                                                                                                                                                   APPLICANT: Liao, Haisun
APPLICANT: Deik, Amy Anderson
APPLICANT: Mamaeva, Natalia
APPLICANT: Woodward, Caroline Ngaara
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEFAX: 619-546-9392
INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                      APPLICANT:
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HYPOTHETICAL: N
ANTI-SENSE: NO
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MEDIUM TYPE: Ploppy disk
COMPUTER: IBM PC compatible
COMPUTER: PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION DATA:
APPLICATION UMBER: US/08/117,952
FILING DATE: 07-SEP-1993
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      REGISTRATION NUMBER: 31,192
REFERENCE/DOCKET NUMBER: P4
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US 08
PILING DATE: 15-JUN-1993
ATTORNEY/AGENT INFORMATION:
NAME: Reiter, Stephen E.
REGISTRATION NUMBER: 31,1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
ADDRESSEE: Pretty, Schroeder, Brueggemann & Clark
STREET: 444 South Flower Street, Suite 2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION:
TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Match 61.1%; Score 11;
Local Similarity 100.0%; Pred. No.
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                                                                                                                                                                                                                                                                                                                                                             INFORMATION:
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RESULT 14
US-08-538-875-24/c
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US-10-341-199-18
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
                            TELEFAX:
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette 3.5inch
                                                                                                                                        APPLICATION NUMBER: KR 93-1751 FILING DATE: 9-FEB-1993 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                      CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE:
                                                                                                                                                                                                                                                                                   OPERATING SYSTEM: MS-DOS
SOPTWARE: WordPerfect 5.1
CURRENT APPLICATION DATA:
              SEQUENCE CHARACTERISTICS:
                                                                          REGISTRATION NUMBER:
REFERENCE/DOCKET NUMBER:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQUENCES: 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                   STREET: #
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STREET: 11/2,
CITY: Incheon
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STREET: #-
STREET: #-
                                                                                                                                                                                                                                                     APPLICATION NUMBER: FILING DATE:
                                                             TELEPHONE:
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                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY: Republic of Korea ZIP: 120-160
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADDRESSEB: Shin, Hang-Cheol
STREET: Jukong Gocheung Apt. 1014-806, Haan-dong
CITY: Kwangmyung-shi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COUNTRY: Republic of Korea
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY: Republic of Korea
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5. 5773582
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#84-4 Daeshin-dong,
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#181-404 Sadang-4-dong, Dongjak-ku
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Shin, Nam-Kyu
Lee, Inkyung
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./2, #302-39 Juan-4-dong, Nam-ku
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100.0%; Pred. No. 3.8e+03;
tive 0; Mismatches 0;
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US-09-142-355B-23/c

US-09-142-355B-23/c

i Sequence 23, Application US/0914235B

i Patent NO. 6184012

i GENERAL INFORMATION:

APPLICANT: Neri, Dario

APPLICANT: Demartie, Salvatore

APPLICANT: Demartie, Salvatore

APPLICANT: Winter, Adrain

APPLICANT: Winter, Gregory Paul

TITLE OF INVENTION: Isolation of Enzymes

FILE REFERENCE: 2224/0E65

CURRENT APPLICATION NUMBER: US/09/142,355B

CURRENT FILING DATE: 1996-09-04

PRIOR APPLICATION NUMBER: GB 9608540.2

PRIOR APPLICATION NUMBER: PCT/GB97/01153

PRIOR PILING DATE: 1997-04-25

PRIOR FILING DATE: 1997-04-25

INUMBER OF SEQ ID NOS: 24

SOPTWARE: FastSEQ for Windows Version 3.0

SEQ ID NO 23

LENGTH: 25

TYPE: DNA
ORGANISM: Artificial Sequence
PEATURE:

OTHER INFORMATION: primer

US-09-142-355B-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; TYPE: nucleic acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: primer DNA
US-08-538-875-24
Search completed: April 7, 2006, 17:01:17 Job time : 106.568 secs
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Maximum DB seq length: 30 /
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Perfect score:
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1: gb ba:*
2: gb-in:*
3: gb-env:*
4: gb-om:*
5: gb-pat:*
7: gb-ph:*
9: gb-pr:*
9: gb-pr:*
10: g6 sts:*
11: g6-sts:*
11: gb-sy:*
13: gb-ui:*
14: gb-htg:*
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18
1 cccccctcttctggatac 18
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Copyright (c) 1993 - 2006 Biocceleration Ltd.
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Regult No.	Score	Query Match	Length	DB	ID	Description	ion
	18	100.0	18	ه إ	CQ796912	CQ796912	Sequence
N	18	100.0	18	σ	CS089158	CS089158	Sequence
ω	18	100.0	18	σ	AX316096	AX316096	Sequence
4	14	77.8	18	0	CS089171	CS089171	Sequence
ۍ	14	77.8	18	σ	AX316109	AX316109	Sequence
6	12	66.7	17	თ	AR568743	AR568743	Sequence
7	12	66.7	17	σ	AX727540	AX727540	Sequence
æ	12	66.7	17	σ	AX733930	AX733930	Sequence
9	12	66,7	17	σ	AX760134	AX760134	Sequence
_	12	66'.7	18	σ	CQ796913	CQ796913	Sequence
c 11	12	66.7	18	σ	CS089159	CS089159	Sequence
c 12	12	66.7	18	σ	AX316097	AX316097	Sequence
13	12	66.7	18	æ	HSRETPO11	х79751 Н.	
C 14	12	66.7	19	თ	AX112294	AX112294	Sequence
c 15	12	66.7	20	σ	AR029828	AR029828	Sequence
16	12	66.7	20	σ	AR311956	AR311956	Sequence
c 17	12	66.7	20	σ	AX298879	AX298879	Sequence
c 18	12	66.7	25	σ	AX112272	AX112272	Sequence

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AR071893	AX117396	AR267420	AR014476	A67134 S	CQ888258	AR634906	AR067084	CS088202	CS064811	AX826876	AR309276	AR305172	BD247806	BD106083	AR144290	AR136351	AX299005	BD169334	BD169223	BD102163	BD102009	AR130022	AX726733	AX724707	AX723024	
Sequence	Sequence	Sequence	Sequence	A67134 Sequence 23	Sequence	Antisense	Novel LDL	Sequence	Sequence	Sequence	Small rem	Degraded	Apoptosis	Agonist a	Sequence	Sequence	Sequence	Sequence								

ALIGNMENTS

RESULT 1

RESULT 2 CSO89158 LOCUS DEFINITION ACCESSION VERSION KEYMORDS SOURCE	р <i>Q</i>	Query Match Best Local : Matches 1:	ORIGIN	FEATURES Bource	JOURNAL	AUTHORS	SOURCE ORGANISM	VERSION KEYWORDS	DEFINITION	CQ796912
CS089158 18 bp DNA Sequence 5 from Patent EP1502948. CS089158 CS089158.1 GI:66714442 . synthetic construct	1 CCCCCTCTTCTGGATAC 18 1 CCCCCTCTTCTGGATAC 18	Query Match 100.0%; Score 18; DB 6; Best Local Similarity 100.0%; Pred. No. 10; Matches 18; Conservative 0; Mismatches		Aquaria Inc. (US) Location/Qualifiers 118	WO 2004026772-A 5 01-APR-2004;	1 Hovanec, T.A. Ammonia-oxidizing bacteria and methods	<pre>synthetic construct synthetic construct other sequences; artificial sequences.</pre>	CQ796912.1 GI:46408538	Sequence 5 from Patent WO2004026772 CQ796912	CO796912 18 bp DNA
linear		; Length 18; 0; Indels	ţ		i	of using and				linear
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25-MAY-2005		Gaps			ı	detecting				PAT 19-APR-2004
2005		0;								2004

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REFERENCE
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Sequence 5 from Patent WO0190312.
                                Hovanec, T.A. and Burrel, P.C.
Ammonia-oxidizing bacteria
Patent: EP 1502948-A 18 02-FEB-2005;
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Patent: EP 1502948-A 5 02-PEB-2005;
Aquaria Inc. (US)
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Patent: WO 0190312-A 5 29-NOV-2001;
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ilarity 100.0%;
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/mol type="unassigned DNA"
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Patent: WO 0190312-A 18 29-NOV-2001;
AQUARIA, INC. (US)
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Unclassified.

1 (bases 1 to 17)

1 (bases 1 to 17)

Gianturco, S.H. and Bradley, W.A.

DNA encoding human apoB48R: a monocyte-macrophage apolipoprotein

B48 receptor gene and protein

B48 receptor gene and protein

Patent: US 6740735-A 25-MAY-2004;
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llarity 100.0%; Pred. No. 2.1e+03;
Conservative 0; Mismatches 0;
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/mol_type="unassigned DNA"
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/note="Description of Artificial Sequence:
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/db_xref="taxon:32630"
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100.0%; Pred. No. 2.9e+04;
tive 0; Mismatches 0;
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Sequences involved in phenomena of tumour suppression, tumour sersion, apoptosis and/or virus resistance and their use as
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Sequence 5564 from Pater
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AX733930.1 GI:30513273
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Sequences involved in phenomena of tumour suppression, tumour suppression, apoptosis and/or virus resistance and their use
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Molecular Engines Laboratories (FR)
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Homo sapiens
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Location/Qualifiers
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                                                                                                                                       /organism="Homo sapiens"
/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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/mol_type="unassigned DNA"
/db_xref="taxon:10090"
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Pred. No. 2.9e+04;
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Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini;
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/db_xref="taxon:32630"
/note="PCR primer"
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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tive 0; Mismatches 0;
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Homo sapiens (human)
Homo sapiens
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Patent: WO 0190312-A 6 29-NOV-2001;
AQUARIA INC. (US)
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Sequence 6 from Patent W00190312.
AX316097.1 GI:17899288
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          Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini; Hominidae; Homo.
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Patent: EP 1502948-A 6 02-FEB-2005;
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/db_xref="taxon:32630"
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/mal_type="unassigned DNA"
/mb.type="unassigned DNA"
/mb.xref="taxon:33630"
/note="Description of Artificial Sequence:
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                                                            Conservative
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                                                                                                                         SEQUENCE"
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Kupperschmitt, A.
Lentiviral triplex dna, and vectors and re
containing lentiviral triplex dna
Patent: WO 0127304-A 23 19-ARR-2001;
INSTITUT PASTEUR (FR) INSTITUT NATIONAL
RECHERCHE MEDICALE (INSEM) (FR)
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mulligan,L.M., Eng,C., Attie,T., Lyonnet,S., Marsh,D.J., Hyland,V.J., Robinson,B.G., Frilling,A., Verellen-Dumoulin,C., Safar,A., Venter,D.J., Munnich,A. and Ponder,B.A.J.

Diverse phenotypes associated with exon 10 mutations of the RET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Direct Submission
Submitted (14-JUN-1994) C. Eng, University of Cambridge, Dept of Pathology, Tennis Court Road, Cambridge CB2 1QP, UK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AX112294.1 GI:13939098
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                                                                                                                                                                                                                                                                                                                                                                                                other sequences; artificial sequences.
                                                                                                                                           /organism="synthetic construct"
/mol_type="unassigned DNA"
/db_xref="taxon:32630"
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/mol_type="genomic DNA"
/isolate="CR3"
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/note="3' end"
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/chromosome="10"
/map="q11.2"
                                                                                                                          'note="MUTATION INTRODUCED INTO THE HIV-1 cPPT CODING
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ocation/Qualifiers
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66.7%; 5-
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100.0%; Pred. No. 2:
tive 0; Mismatches
        Score 12; DB; Pred. No. 2:
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Maximum Match 100%
Listing first 45 summaries
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     Published Applications NA New:*

1: #./SIDS5/ptodate/1/pubpna/US06_NEW_PUB.seq:*

2: ** /SIDS5/ptodata/1/pubpna/US06_NEW_PUB.seq:*

3: /SIDS5/ptodata/1/pubpna/US07_NEW_PUB.seq:*

4: /SIDS5/ptodata/1/pubpna/US07_NEW_PUB.seq:*

4: /SIDS5/ptodata/1/pubpna/US09_NEW_PUB.seq:*

6: /SIDS5/ptodata/1/pubpna/US10_NEW_PUB.seq:*

7: /SIDS5/ptodata/1/pubpna/US10_NEW_PUB.seq:*

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14: /SIDS5/ptodata/1/pubpna/US11_NEW_PUB.seq:*

15: /SIDS5/ptodata/1/pubpna/US11_NEW_PUB.seq:*
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     GenCore version 5.1.7 (c) 1993 - 2006 Biocceleration Ltd.
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          Sequence 112898,
Sequence 1578545,
Sequence 1578545,
Sequence 352597,
Sequence 352597,
Sequence 352632,
Sequence 352632,
Sequence 84927, A
Sequence 84927, A
Sequence 1353078,
Sequence 1473850,
Sequence 1473850,
Sequence 1473850,
Sequence 1173850,
Sequence 1173850,
Sequence 1173850,
Sequence 1899, A
Sequence 1896978,
Sequence 1896978,
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σ		Sequence 447881, Sequence 1261081, Sequence 466300, Sequence 899298, Sequence 1261104.	Sequence 96730, Sequence 96731, Ap Sequence 36513, Ap Sequence 171065, Sequence 171065, Sequence 278911 Sequence 1230837, Sequence 1230837, Sequence 224598, Sequence 224598, Sequence 8320, A Sequence 836977, Sequence 896977, Sequence 896977, Sequence 2344, App	

ALIGNMENTS

RESULT 1
US-10-310-914A-1061199/c
US-10-310-914A-1061199/c
; Sequence 1061199, Application US/10310914A
; Publication No. US20060003322A1
; GENERAL INFORMATION:

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APPLICANT: Bentwich, Isaac
APPLICANT: Shiler, Kvuzat
TITLE OF INVENTION: Bioinformatically detectable group of novel regulatory genes an
ITITLE OF INVENTION: uses thereof
FILE REFERENCE: 06087.0200.CPUS01
CURRENT APPLICATION NUMBER: US/10/310,914A
CURRENT FILING DATE: 2002-12-06
NUMBER OF SEQ ID NOS: 1388402
SOFTWARE: PatentIn version 3.3
SEQ ID NO 1061199
LENGTH: 25
TYPE: RNA
ORGANISM: Human
US-10-310-914A-1061199
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US-11-121-849-112898
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Sequence 112899, Application US/11121849
Publication No. US20050272080A1
GENERAL INFORMATION:
APPLICANT: John Palma
TITLE OF INVENTION: Methods of Genetic Analysis of Formalin Fixed Paraffin Embedded
TITLE OF INVENTION: Microarrays
FILE REFERENCE: 3684.1
CURRENT APPLICATION NUMBER: US/11/121,849
CURRENT FILING DATE: 2005-05-03
PRIOR APPLICATION NUMBER: 60/567,949
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Best Local Similarity 100.
Matches 14; Conservative
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100.0%; Pr
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APPLICANT: Khvorova, Anastasia
APPLICANT: Reynolds, Angela
APPLICANT: Leake, Devin
APPLICANT: Leake, Devin
APPLICANT: Marshall, William
APPLICANT: Scaringe, Stephen
TITLE OF INVENTION: Functional and Hyperfunctional sirna
FILE REFERENCE: 13499US
CURRENT APPLICATION NUMBER: US/11/101,244
CURRENT APPLICATION NUMBER: 00/5-04-07
FRIOR APPLICATION NUMBER: 60/50,050
PRIOR TILING DATE: 2003-09-10
PRIOR APPLICATION NUMBER: 60/426,137
PRIOR APPLICATION NUMBER: 60/426,137
PRIOR PILING DATE: 2002-11-14
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                                                                                                                                                                                                                                                                                                                                                                                         US-11-083-784-1578545
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US-11-101-244-1578545
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NUMBER OF SEQ ID NOS: 673904
SOFTWARE: Microarray Probe Sequence Listing Generator V 1.1
SEQ ID NO 112898
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Matches
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SEQ ID NO 1578545
LENGTH: 19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
                                                                                                                                                 APPLICANT: Dharmacon, Inc.
APPLICANT: Khvorova, Anastasia
APPLICANT: Reymolds, Angela
APPLICANT: Leake, Devin
APPLICANT: Marshall, William
APPLICANT: Scaringe, Stephen
APPLICANT: Scaringe, Stephen
APPLICANT: Scaringe, Stephen
CURRENT APPLICATION NUMBER: US/11/083,784
CURRENT FILING DATE: 2005-03-18
PRIOR APPLICATION NUMBER: US/10/714,333
PRIOR FILING DATE: 2003-11-14
PRIOR APPLICATION NUMBER: 60/502,050
PRIOR FILING DATE: 2003-09-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Dharmacon, Inc. APPLICANT: Khvorova, Anas
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                                                                                                                                       FILE REFERENCE: 13499US
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es 14; Conservative 0;
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Similarity 64.7%; Pred. No. 8.5e+02;
ll; Conservative 4; Mismatches 2
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; TYPE: DNA; ORGANISM: Homo sapien US-11-121-849-325988
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US-11-083-784-1578545
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US-11-136-527-352597, Application US/11136527
; Sequence 352597, Application US/11136527
; Publication No. US20050287570A1
; GENERAL INFORMATION:
                                     ; OTHER INFORMATION: Probe US-11-136-527-352597
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Publication No. US20050272080A1
GENERAL INFORMATION:
APPLICANT: John Palma
TITLE OF INVENTION: Methods of Genetic Analysis of Formalin Fixed Paraffin Embedded
TITLE OF INVENTION: Microarrays
FILE REFERENCE: 3684.1
CURRENT APPLICATION NUMBER: US/11/121,849
CURRENT FILING DATE: 2005-05-03
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SOFTWARE: Microarray Probe Sequence Listing Generator V 1.1
SEQ ID NO 325988
LENGTH: 25
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PRIOR FILING DATE: 2002-11-14
NUMBER OF SEQ ID NOS: 1591911
SOFTWARE: Proprietary
SEQ ID NO 1578545
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Best Local Similarity
Matches 15; Conserv
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Best Local Similarity
                                                                                                                                                         SEQ ID NO 352597
                                                                                                                                                                                                                                                          APPLICANT: Wyeth
APPLICANT: Mounts, William M
APPLICANT: Mounts, William M
TITLE OF INVENTION: Probe Arrays For Expression Profiling of Rat Genes
FILE REFERENCE: 031896-041000 (AMI01086)
CURRENT APPLICATION NUMBER: US/11/136,527
CURRENT FILING DATE: 2005-05-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT FILING DATE: 2005-05-03
PRIOR APPLICATION NUMBER: 60/567,949
PRIOR FILING DATE: 2004-05-03
                                                                                                                                                                          PRIOR APPLICATION NUMBER: US 60/574,294
PRIOR FILING DATE: 2005-05-26
NUMBER OF SEQ ID NOS: 362830
SOFTWARE: PatentIn version 3.2
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                                                                           FEATURE:
                                                                                            ORGANISM: Artificial
                                                                                                                                      LENGTH: 25
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88.2%;
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64.7%;
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DB 14;
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Length 25;
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Query Match

76.7%;

Score 13.8;

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15;

CCCCCTCTTCTGGATAC 18

Matches

Conservative

88.2%;

Pred. No. 9e+02; 0; Mismatches

<u>ب</u>

Indels

0

Gaps

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Best Local Similarity

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                                                                                                                                  ; OTHER INFORMATION: Probe US-11-136-527-352632
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APPLICANT: Wounts, William M
APPLICANT: Mounts, William M
TITLE OF INVENTION: Probe Arrays For Expression Pro
FILE REFERENCE: 031896-041000 (AM101086)
CURRENT APPLICATION NUMBER: US/11/136,527
CURRENT FILING DATE: 2005-05-25
PRIOR APPLICATION NUMBER: US 60/574,294
PRIOR FILING DATE: 2005-05-26
NUMBER OF SEQ ID NOS: 362830
SOPTWARE: Patentin version 3.2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; OTHER INFORMATION: Probe US-11-136-527-352621
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US-11-136-527-352621/c
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SEQ ID NO 352632
LENGTH: 25
                                                                                                Query Match
                                                                                                                                                                                                                                                       APPLICANT: Wyeth
APPLICANT: Mounts, William M
APPLICANT: Mounts, William M
TITLE OF INVENTION: Probe Arrays For Expression Profiling of Rat Genes
FILE REFERENCE: 031996-041000 (AM101086)
CURRENT APPLICATION NUMBER: US/11/136,527
CURRENT FILING DATE: 2005-05-25
PRIOR APPLICATION NUMBER: US 60/574,294
PRIOR FILING DATE: 2005-05-26
NUMBER OF SEQ ID NOS: 362830
COMMUNICATION NUMBER: 362830
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Best Local Similarity
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                                                                                                                                                                FEATURE:
                                                                                                                                                                            ORGANISM: Artificial
                                                                                                                                                                                                    TYPE: DNA
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                                                                             Similarity
                             CCCCCTCTTCTGGATAC 18
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88.2%;
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                                                           Score 13.8; DB Pred. No. 9e+02; O; Mismatches
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RESULT 9

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US-10-310-914A-84927
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APPLICANT: Shiler, Kvuzat
TITLE OF INVENTION: Bioinformatically detectable group of novel regulatory genes
TITLE OF INVENTION: uses thereof
TITLE OF INVENTION: Uses thereof
TITLE REFERENCE: 06087.0200.CPUS01
CURRENT APPLICATION NUMBER: US/10/310,914A
CURRENT APPLICATION NUMBER: US/10/310,914A
CURRENT PILING DATE: 2002-12-06
NUMBER OF SEQ ID NOS: 1388402
SOFTWARE: Patentin version 3.3
SEQ ID NO 84927
APPLICANT: John Palma
TITLE OF INVENTION: Methods of Genetic Analysis of Formalin Fixed Paraffin Embedded
TITLE OF INVENTION: Microarrays
FILE REFERENCE: 3684.1
CURRENT APPLICATION NUMBER: US/11/121,849
CURRENT FILING DATE: 2005-05-03
PRIOR APPLICATION NUMBER: 60/567,949
PRIOR FILING DATE: 2004-05-03
NUMBER OF SEQ ID NOS: 673904
SOFTMARE: Microarray Probe Sequence Listing Generator V 1.1
SEQ ID NO 394245
                                                                                                                                                                                                                                 GENERAL INFORMATION:
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APPLICANT: Shiler, Kvuzat
TITLE OF INVENTION: Bioinformatically detectable group of novel regulatory genes
TITLE OF INVENTION: uses thereof
FILE REFERENCE: 06087.0200.CPUS01
CURRENT FILING DATE: 2002-12-06
NUMBER OF SEQ ID NOS: 1388402
SOFTWARE: PatentIn version 3.3
SEQ ID NO 908031
                                                                                                                                                                                                                                                       Sequence 394245, Application US/11121849 Publication No. US20050272080A1
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Best Local :
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73.38;
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93.3%;
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Pred. No. 1.5e+03;
3; Mismatches 1
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APPLICANT: Khvorova, Anastasia
APPLICANT: Reynolds, Angela
APPLICANT: Leake, Devin
APPLICANT: Leake, Devin
APPLICANT: Marshall, William
APPLICANT: Scaringe, Stephen
TITLE OF INVENTION: Functional and Hyperfunctional sirNA
FILE REFERENCE: 13499US
CURRENT APPLICATION NUMBER: US/11/101,244
CURRENT APPLICATION NUMBER: 007
PRIOR APPLICATION NUMBER: 60/502,050
PRIOR FILLING DATE: 2003-09-10
PRIOR APPLICATION NUMBER: 60/426,137
PRIOR APPLICATION NUMBER: 60/426,137
PRIOR FILLING DATE: 2002-11-14
PRIOR FILLING DATE: 2002-11-14
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US-10-310-914A-1353078/c
US-10-310-914A-1353078, Application US/10310914A
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; TYPE: DNA
; ORGANISM: Homo sapien
US-11-121-849-394245
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                                                                         ; ORGANISM: Homo sapiens
US-11-101-244-1473850
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APPLICANT: Shiler, Kvuzat
APPLICANT: Shiler, Kvuzat
TITLE OF INVENTION: Bioinformatically detectable group of novel regulatory genes and
TITLE OF INVENTION: uses thereof
FILE REFERENCE: 06087.0200.CPUS01
CURRENT APPLICATION NUMBER: US/10/310,914A
CURRENT FILING DATE: 2002-12-06
NUMBER OF SEQ ID NOS: 1388402
SOFTWARE: Patentin version 3.3
SEQ ID NO 1353078
LENGTH: 19
TWORD: DATA
Query Match 73.3%;
Best Local Similarity 61.1%;
Matches 11; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION: APPLICANT: Dharmacon,
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Publication No. US20050246794A1
GENERAL INFORMATION:
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                                                                                                                             SOFTWARE: Proprietary
SEQ ID NO 1473850
LENGTH: 19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ORGANISM: Human
-10-310-914A-1353078
                                                                                                                                                                                  NUMBER OF SEQ ID NOS: 1591911
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: RNA
                                                                                                               TYPE: RNA
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93.3%;
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83.3%; Pred. No. 1.
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Pred. No. 1.5e+03;
0; Mismatches 1;
 Score 13.2; DB 12
Pred. No. 1.8e+03;
4; Mismatches 3
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                                    DB 12;
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US-11-083-784-1473850
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US-10-770-726-11899/c
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US-11-083-784-1473850
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US-10-770-726-11899
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CURRENT FILING DATE: 2005-03-18
PRIOR APPLICATION NUMBER: US/10/714,333
PRIOR FILING DATE: 2003-11-14
PRIOR APPLICATION NUMBER: 60/502,050
PRIOR APPLICATION NUMBER: 60/426,137
PRIOR APPLICATION NUMBER: 60/426,137
PRIOR FILING DATE: 2002-11-14
PRIOR FILING DATE: 2002-11-14
                                                                                                                                                                                                         APPLICANT: Wyeth
APPLICANT: Brown, Eugene
APPLICANT: Liu, Mei
APPLICANT: Liu, Mei
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR DIAGNOSING, PREVENTING, AND TREATI?
TITLE OF INVENTION: CANCERS
FILE REPERENCE: AM101079 (031896-010000)
CURRENT APPLICATION NUMBER: US/10/770,726
CURRENT FILING DATE: 2004-02-04
NUMBER OF SEQ ID NOS: 48640
SOFTWARE: PATENTIN VETSION 3.2
SEQ ID NO 11899
                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 11899, Application US/10770726 Publication No. US20050266409A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: Proprietary
SEQ ID NO 1473850
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Best Local :
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APPLICANT: Scaringe, Stephen
TITLE OF INVENTION: Functional and Hyperfunctional siRNA
FILE REFERENCE: 13499US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQ ID NOS: 1591911
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                                                                                                                                                                         LENGTH: 21
TYPE: RNA
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                                                                                    Local Similarity
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Local Similarity 61.1%;
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Reynolds, Angela
Leake, Devin
                                                                      Conservative
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                                                                                      Score 13.2;
Pred. No. 1.
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GenCore version 5.1.7
Copyright (c) 1993 - 2006 Biocceleration Ltd.

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1: /cgn2_6/ptodata/1/pubpna/US07_PUBCOMB.seq:*

2: /cgn2_6/ptodata/1/pubpna/US08_PUBCOMB.seq:*

3: /cgn2_6/ptodata/1/pubpna/US08_PUBCOMB.seq:*

4: /cgn2_6/ptodata/1/pubpna/US08_PUBCOMB.seq:*

5: /cgn2_6/ptodata/1/pubpna/US108_PUBCOMB.seq:*

6: /cgn2_6/ptodata/1/pubpna/US108_PUBCOMB.seq:*

7: /cgn2_6/ptodata/1/pubpna/US108_PUBCOMB.seq:*

9: /cgn2_6/ptodata/1/pubpna/US108_PUBCOMB.seq:*

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        7 US-10-659-948A-5

7 US-10-659-980A-5

7 US-10-719-956-558629

9 US-09-320-337-74

9 US-10-659-948A-21

10 US-10-659-948A-21

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10 US-10-19-956-50560

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10 US-10-19-956-470164

10 US-10-719-956-470164

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10 US-11-060-756-206014

10 US-11-060-756-206016

10 US-10-719-900-48801

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10 US-10-719-900-680676

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10 US-10-719-900-680676

10 US-10-719-900-680676

10 US-10-719-900-6800
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Sequence 5, Appli
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Sequence 71, Appl
Sequence 21, Appl
Sequence 21, Appl
Sequence 21, Appl
Sequence 2535, A
Sequence 50560, A
Sequence 94286, A
Sequence 94286, A
Sequence 94286, A
Sequence 94281, Appl
Sequence 338352,
Sequence 416880,
Sequence 202014,
Sequence 288113, A
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US-10-809-189-56205	US-10-719-900-949533	US-10-719-900-904924	US-10-719-900-761202	US-10-719-900-627829	US-10-719-900-550905	US-10-719-900-308249	US-10-719-900-226025	US-10-719-900-28087	US-10-719-956-558628	US-10-719-956-89806	US-10-098-263B-37037	US-10-459-876-17	US-11-060-756-197598	US-11-060-756-197597	US-10-719-900-574815	US-10-719-900-316050	US-10-719-956-611802	US-10-719-956-511003	US-10-719-956-193006	US-10-719-956-117959	US-10-282-122A-78603
Sequence	Sequence	Sequence	Sequence	Sequence	Sequence		Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence
56205. A	949533,	904924,	761202,	627829,	550905,	308249,	226025,	28087, A	558628,	89806, A	37037, A	17, Appl	Sequence 197598,	Sequence 197597,	574815,	316050,	611802,	511003,	193006,	117959,	78603, A

ALIGNMENTS

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GENERAL INFORMATION:

APPLICANT: HOVANEC, Timothy A

ITITE OF INVENTION: Method of Using Ammonia-Oxidizing Bacteria

FILE REFERENCE: 81289-294309

CURRENT APPLICATION NUMBER: US/10/659,948A

CURRENT FILING DATE: 2003-09-10

PRIOR APPLICATION NUMBER: US 09/573,684

PRIOR FILING DATE: 2000-05-19

PRIOR APPLICATION NUMBER: US 60/386,217

PRIOR APPLICATION NUMBER: US 60/386,217

PRIOR FILING DATE: 2002-09-19

INUMBER OF SEQ ID NOMBER: US 60/386,219

PRIOR APPLICATION NUMBER: US 60/386,219

PRIOR PILING DATE: 2002-09-19

NUMBER OF SEQ ID NOS: 23

SOPTWARE: Patentin version 3.2

SEQ ID NO 5

LENGTH: 18

TYPE: DNA

ORGANISM: Artificial
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                                                                         Sequence 5, Application US/10659980A Publication No. US20040106133A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 5, Application US/10659948A Publication No. US20040101946A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
Matches 18; Conserv
APPLICANT: Hovanec, Timothy A TITLE OF INVENTION: Method for Detecting Ammonia-Oxidizing Bacteria FILE REFERENCE: 81289-284781
                                                                                                                                                                                                                                                                                      1 CCCCCCTCTTCTGGATAC 18
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CURRENT APPLICATION NUMBER: US/10/659,980A

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RESULT 3
US-10-659-983A-5
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US-10-719-956-558629
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; OTHER INFORMATION: Oligonucleotide Probe
US-10-659-983A-5
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PRIOR APPLICATION NUMBER: US 09/573,684
PRIOR ELLING DATE: 2000-05-19
PRIOR APPLICATION NUMBER: US 60/386,217
PRIOR PILING DATE: 2002-09-19
PRIOR APPLICATION NUMBER: US 60/386,218
PRIOR PILING DATE: 2002-09-19
PRIOR PILING DATE: 2002-09-19
PRIOR APPLICATION NUMBER: US 60/386,219
PRIOR PILING DATE: 2002-09-19
                                                            Sequence 558629, Application US/10719956 Publication No. US20040146910A1 GENERAL INFORMATION:
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LENGTH: 18
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SEQ ID NO 5
LENGTH: 18
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APPLICANT: Xue Mei Zhou
TITLE OF INVENTION: Methods of Genetic Analysis of Rat
FILE REFERENCE: 3527.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION NUMBER: US 60/386,217
PRIOR FILING DATE: 2002-09-19
PRIOR APPLICATION NUMBER: US 60/386,218
PRIOR FILING DATE: 2002-09-19
PRIOR APPLICATION NUMBER: US 60/386,219
PRIOR FILING DATE: 2002-09-19
PRIOR FILING DATE: 2002-09-19
NUMBER OF SEQ ID NOS: 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION: Ammonia-Oxidizing Bacteria FILE REFERENCE: 81289-284779 CURRENT APPLICATION NUMBER: US/10/659,983A CURRENT FILING DATE: 2003-09-10
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                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: DNA
ORGANISM: Artificial
                                                                                                                                                                                                                                                                                                            Local Similarity
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100.0%; Pred. No. 36;
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; NUMBER OF SEQ ID NOS: 699466
; SOFTWARE: Microarray Probe Sequence Listing Generator V 1.1
; SEQ ID NO 558629
; LENGTH: 25
; TYPE: DNA
; ORGANISM: Rattus norvegicus
US-10-719-956-558629
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Best Local Similarity
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                                                                                                ; MOLECULE TYPE: oligonucleotide US-09-320-337-74
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Best Local S
Matches 15
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                                                                                                                                                                                                                                                                                                                         CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/442
PILING DATE: 17-MAY-1995
APPLICATION NUMBER: 08/245
PILING DATE: 18-MAY-1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION NUMBER: 60/427,836 PRIOR FILING DATE: 2002 11 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION NUMBER: US/10/719,956
CURRENT FILING DATE: 2003-11-20
                                                                                                                                                                                                              TELEPHONE: 973-994-174
TELEPAX: 973-994-1744
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/0
FILING DATE: 26-MAY-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER: IBM FACTOR OPERATING SYSTEM: WINI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Bohinski, Robert J.,
APPLICANT: Wiltsett, Jeffrey A.
TITLE OF INVENTION: Nucleic Acid
TITLE OF INVENTION: Lung Cell-
                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                REFERENCE/DOCKET NUMBER: 27
TELECOMMUNICATION INFORMATION:
TELEPHONE: 973-994-1700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM: MEDIUM TYPE: 3.5 incl
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                                                                                                                             TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                               NAME: Olstein, Elliot M. REGISTRATION NUMBER: 24,025
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CITY: Roseland
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADDRESSEE:
                                                                                                                                                                                 LENGTH: 30 bases
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15; Conserv
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CCCCCTCTTCTGGATA 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  6 Becker Farm Road
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320010016352A1
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                                  Conservative
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Nucleic Acid Sequences Controlling
Lung Cell - Specific Gene Expression
76
                                                93.8%;
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Pred. No. 2.4e+03;
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                                                 Score 14.4; DB 3;
Pred. No. 2.4e+03;
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                                    Mismatches
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APPLICANT: Hovenec, Timothy A
ITILE OF INVENTION: Method for Detecting Ammonia-Oxidizing Bacteria
FILE REFERENCE: 81289-284781
CURRENT APPLICATION NUMBER: US/10/659,980A
CURRENT FILING DATE: 2003-09-10
PRIOR APPLICATION NUMBER: US 09/573,684
PRIOR FILLING DATE: 2000-05-19
PRIOR APPLICATION NUMBER: US 60/386,217
PRIOR FILING DATE: 2002-09-19
PRIOR APPLICATION NUMBER: US 60/386,218
PRIOR FILING DATE: 2002-09-19
PRIOR APPLICATION NUMBER: US 60/386,219
PRIOR APPLICATION NUMBER: US 60/386,219
PRIOR FILING DATE: 2002-09-19
SOFTWARE: Patentin version 3.2
SOFTWARE: Patentin version 3.2
LENGTH: 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-10-659-980A-21
; Sequence 21, Application US/10659980A
; Publication No. US20040106133A1
; GENERAL INFORMATION:
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                                                                                                            ; OTHER INFORMATION: Oligonucleotide probe US-10-659-980A-21
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CURRENT FILING DATE: 2003-09-10
PRIOR APPLICATION NUMBER: US 09/573,684
PRIOR APPLICATION NUMBER: US 60/386,217
PRIOR APPLICATION NUMBER: US 60/386,217
PRIOR APPLICATION NUMBER: US 60/386,218
PRIOR APPLICATION NUMBER: US 60/386,218
PRIOR PILING DATE: 2002-09-19
PRIOR APPLICATION NUMBER: US 60/386,219
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TITLE OF INVENTION: Method of Using Ammonia-Oxidizing Bacteria
FILE REFERENCE: 81289-294309
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                                                                                                                                                          TYPE: DNA
ORGANISM: Artificial
FEATURE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: DNA
  Y Match 76.7%; Score 13.8; DB 7; Length 18; Local Similarity 88.2%; Pred. No. 4.9e+03; hes 15; Conservative 0; Mismatches 2; Indels
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                                                                                                                                                     ; LENGTH: 25
; TYPE: DNA
; ORGANISM: Rattus norvegicus
US-10-719-956-25535
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PRIOR FILING DATE: 2002-09-19
PRIOR APPLICATION NUMBER: US 60/386,218
PRIOR FILING DATE: 2002-09-19
PRIOR FILING DATE: 2002-09-19
PRIOR APPLICATION NUMBER: US 60/386,219
PRIOR FILING DATE: 2002-09-19
PRIOR FILING DATE: 2002-09-19
NUMBER OF SEQ ID NOS: 23
SOPTWARE: PatentIn version 3.2
                                                                                                                                                                                                                                     SOPTWARE: Microarray Probe Sequence Listing Generator V 1.1 SEQ ID NO 25535
                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 25535, Application US/10719956
Publication No. US20040146910A1
GENERAL INFORMATION:
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Publication No. US20040157313A1
GENERAL INFORMATION:
APPLICANT: Hovanec, Timothy A
TITLE OF INVENTION: Ammonia-Oxidizing Bacteria
FILE REFERENCE: 81289-284779
CURRENT APPLICATION NUMBER: US/10/659,983A
CURRENT FILING DATE: 2003-09-10
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LENGTH: 18
                                                                                Matches
                                                                                               Query Match
Best Local (
                                                                                                                                                                                                                                                            APPLICANT: Xue Mei Zhou
TITLE OF INVENTION: Methods of Genetic Analysis of Rat
FILE REFERENCE: 3527.1
CURRENT APPLICATION NUMBER: US/10/719,956
CURRENT FILING DATE: 2003-11-20
PRIOR APPLICATION NUMBER: 60/427,836
PRIOR APPLICATION NUMBER: 60/427,836
PRIOR FILING DATE: 2002 11 20
NUMBER OF SEQ ID NOS: 699466
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION NUMBER: US 09/573,684 PRIOR FILING DATE: 2000-05-19
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ORGANISM: Artificial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FEATURE:
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                          1 CCCCCCTCTTCTGGATA 17
                                                                                                 Similarity
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CCGCCCTGTTCTGGATA 19
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                                                                              Conservative
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88.2%;
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88.2%;
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Pred. No. 4.9e+03;
                                                                          Score 13.8; DB 7;
Pred. No. 4.8e+03;
0; Mismatches 2
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                                                                                                               DB 7; Length 25;
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RESULT 10 US-10-719-956-50560/c

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; Sequence 50560, Application US/10719956
; Publication No. US20040146910A1
; GENERAL INFORMATION:
APPLICANT: Xue Mei Zhou
; TITLE OF INVENTION: Methods of Genetic Analysis of Rat
; FILE REFERENCE: 3527.1
; CURRENT APPLICATION NUMBER: US/10/719,956
; CURRENT FILING DATE: 2003-11-20
; PRIOR APPLICATION NUMBER: 60/427,836
; PRIOR APPLICATION NUMBER: 60/427,836
; PRIOR FILING DATE: 2002 11 20
; NUMBER OF SEQ ID NOS: 659466
; SOFTMARE: Microarray Probe Sequence Listing Generator V 1.1
; SEQ ID NO 50560
Sequence 94288, Application US/10719956

Publication No. US20040146910A1

GENERAL INFORMATION:
APPLICANT: XUE Mei Zhou
TITLE OF INVENTION: Methods of Genetic Analysis of TILE REFERENCE: 3527.1

CURRENT PILING DATE: 2003-11-20
PRIOR APPLICATION NUMBER: 60/427,836

PRIOR PILING DATE: 2002 11 20
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APPLICANI: Goetting, Christian
APPLICANI: Goetting, Christian
APPLICANI: Goetting, Christian
APPLICANI: Goetting, Christian
APPLICANI: Kuhn, Joachim
ITILE OF INVENTION: Xylosyltransferase and isc
FILE REFERENCE: 020195-00002
CURRENT APPLICATION NUMBER: US/10/169,368
CURRENT FILING DATE: 2002-07-01
PRIOR APPLICATION NUMBER: PCT/EP00/13311
PRIOR APPLICATION NUMBER: PCT/EP00/13311
PRIOR FILING DATE: 2000-12-30
NUMBER OF SEQ ID NOS: 36
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 13
LENGTH: 28
LENGTH: 28
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US-10-719-956-94288
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; ORGANISM: Rattue norvegicue
US-10-719-956-50560
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; OTHER INFORMATION: Description of Artificial Sequence: US-10-169-368-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-10-169-368-13/c
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Best Local S
Matches 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FEATURE:
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Similarity 88.2%; Pred. No. 4.8e+03;
15; Conservative 0; Mismatches 2
                                                                                                                                                                                                                                                                                                               CCCCACTTCTGCATAC 5
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88.2%;
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Pred. No. 4.8e+03;
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; LENGTH: 25
; TYPE: DIA
; ORGANISM: Rattus norvegicus
US-10-719-956-94288
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; Sequence 338352, Application US/10719900
; Publication No. US20050026164A1
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; TYPB: DNA
; ORGANISM: Rattus norvegicus
US-10-719-956-470164
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                                                                                                                                    ; ORGANISM: Mus musculus US-10-719-900-338352
                                                                                                                                                                                                GENERAL INFORMATION:

APPLICANT: Xue Mei Zhou
APPLICANT: Xue Mei Zhou
TITLE OF INVENTION: Methods of Genetic Analysis of Mouse
TITLE OF INVENTION: Methods of Genetic Analysis of Mouse
TILE REFERENCE: 3528.1

CURRENT FILING DATB: 2003-11-20
PRIOR APPLICATION NUMBER: 60/427,808
PRIOR FILING DATB: 2002 11 20
NUMBER OF SEQ ID NOS: 982914

SOFTWARE: Microarray Probe Sequence Listing Generator V 1
SEQ ID NO 338352
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Xue Mei Zhou
TITLE OF INVENTION: Methods of Genetic Analysis of Rat
FILE REFERENCE: 3527.1
CURRENT APPLICATION NUMBER: US/10/719,956
CURRENT FILING DATE: 2003-11-20
PRIOR APPLICATION NUMBER: 60/427,836
PRIOR FILING DATE: 2002 11 20
NUMBER OF SEQ ID NOS: 699466
SOFTWARE: Microarray Probe Sequence Listing Generator V 1.1
SEQ ID NO 470164
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 470164, Application US/10719956 Publication No. US20040146910A1 GENERAL INFORMATION:
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SOFTWARE: Microarray Probe Se
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l Similarity 93.3%;
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Pred. No. 7.7e+03;
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US-10-719-900-416880

US-10-719-900-416880, Application US/10719900

Sequence 41680, Application US/10719900

Publication No. US20050026164A1

GENERAL INFORMATION:

APPLICANT: Xue Med EDOU

TITLE OF INVENTION: Methods of Genetic Analysis of Mouse

FILE REFERENCE: 3528.1

CURRENT FILING DATE: 2003-11-20

PRIOR APPLICATION NUMBER: 60/427,808

PRIOR APPLICATION NUMBER: 60/427,808

PRIOR APPLICATION NUMBER: 60/427,808

PRIOR PILING DATE: 2002-11-20

PRIOR APPLICATION NUMBER: 60/427,808

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PRIOR APPLICATION NUMBER: 60/427,808

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                   Sequence 74, Appl Sequence 17, Appl Sequence 17, Appl Sequence 2493, Ap Sequence 305, Ap Sequence 24539, A Sequence 277, Appl Sequence 27092, A Sequence 270, Appl Sequence 28, Appl Sequence 28, Appl Sequence 28, Appl Sequence 28, Appl Sequence 17, Appl Sequence 17, Appl Sequence 17, Appl Sequence 11, Appl Sequence 16, Appl Sequence 10, Appl Sequence 3, Appl Sequence 290, App Sequence 290, App Sequence 290, App Sequence 290, App Sequence 290, App
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63.3	63.3	63.3	63.3	64.4	64.4	64.4	64.4	64.4	64.4	64.4	64.4	64.4	64.4	64.4	64.4	64.4	64.4	64.4	64.4	64.4
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			5423	42	-	Sequence 1, Appli	ŗ			Sequence 103966,		Seguence 10214, A	Sequence 11872, A	Sequence 11871, A	Sequence 11870, A	11869,	Sequence 11868, A	11867,	Sequence 11866, A	Sequence 11865, A

ALIGNMENTS

RESULT 1

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INFORMATION FO SEQUENCE CHA SEQUENCE CHA LENGTH: 3 TYPE: NUC STRANDEDNE FOPOLOGY: MOLECULE TYP US-08-442-809A-74	TELLIA	# G	_8	US-08-442-809A-74/c ; Sequence 74, Appli ; Patent No. 5976873 ; GENERAL INFORMATI: Boh APPLICANT: Boh APPLICANT: Whi TITLE OF INVENT ; TITLE OF INVENT ; TITLE OF SEQUE ; CORRESPONDENCE
TELEF RMATI QUENC LENGT TYPE: STRAN TOPOL LECUL	APPLI FILIN TORNE NAME: REGIS REFER TRLECOM	PIL	ADDRE ADDRE STREE CITY: CITY: STATE COUNT COUNT COUNT EDITE	-442-80
TELEFAX: FORMATION SEQUENCE CI SEQUENCE CI LENGTH: TYPE: IN STRANDEDI TOPOLOGY MOLECULE T: -442-809A-*	ERE ING	PUI RAI TWA INC SSI	ADDRESSE ADDRESSE ADDRESSE STREET: CCITY: R STATE: STATE: O7 ZIP: 07 MEDIUM T	-442-809A- uence 74, ent No. 59 NERAL INFO APPLICANT: APPLICANT: TITLE OF I TITLE OF I TITLE OF I TITLE OF I
TELEFAX: RMATION PO QUENCE CH; LENGTH: TYPE: TYPE: TOPOLOGY: LECULE TYE LECULE TYE LECULE TYE	APPLICATIO FILING DAT TORNEY/AGE NAMB: Ols REGISTRATI REFERENCE/ LECOMMUNIC	COMPUTER: COMPUTER: COMPUTER: SOPTWARE: SOPTWARE: RRENT APP RRENT APP APPLICATI FILING DA CLASSIFIC IOR APPLI	ADDRESSEE: ADDRESSEE: STREET: 6 STREET: ROS CITY: ROS COUNTE: NO COUNTE: NO ZIP: 0706 MPUTER REA MPUTER REA MPUTER REA	NT:
N K KOLIS KI		COMPUTER: I OPERATING SY SOFTWARE: W ROFTWARE: W RRENT APPLICATION: APPLICATION DATE: FILING DATE: CLASSIFICATIO IOR APPLICATI	ADDRESSEE: ADDRESSEE: STREET: 6 B CITY: Rosel STATE: New COUNTRY: US ZIP: 07068 MPUTER READA MPUTER READA	RAPA RAPA RAPA RAPA RAPA RAPA RAPA RAPA
1 1. 0.0 1	APPLICATION NUMBER: 08/2 FILING DATE: 18-MAY-1994 ATTORNEY/AGENT INFORMATION: NAME: Olstein, Elliot M.; REGISTRATION NUMBER: 24, REFERENCE/DOCKET NUMBER: TELECOMMUNICATION INFORMATI TELECHONE: 201-204-1700	HO ZEOM	ADDRESSEE: Carell ADDRESSES: Cecchi STREET: 6 Becker CITY: Roseland STATE: New Jersey COUNTRY: USA ZIP: 07068 MEDIUM TYPR: 1 5	08-442-809A-74/c equence 74, Application US/08442809A atent No. 5976873 GENERAL INFORMATION: APPLICANT: Bohinski, Robert J., APPLICANT: Whitsett, Jeffrey A. TITLE OF INVENTION: Nucleic Acid TITLE OF INVENTION: Controlling L TITLE OF INVENTION: Specific Gene NUMBER OF SEQUENCES: 76 CORRESPONDENCES: 76
SEQ I SEQ I CTERI bases ic ac inear		MBI 17	Carel Cecch Cecker and Jerse A	ON SECOND
994-174 90 ID NO: 74: ERISTICS: ses acid single single oligonucleotide	ER: 08/245, -MAY-1994 ORMATION: Elliot M. BER: 24,025 NUMBER: 27 INFORMATION:	M FS/2 TEM: MS-DOS TAPERFECT 5. TION DATA: TIOMBER: US/0 17-MAY-1995 ON DATA:	Carella, Carella, Cecchi, Secker Fai and Jersey A BLE FORM:	ki, tt,
7444 ICS ICS	PATION OF THE PARTY OF THE PART	MS-DOS ect 5. ATA: US/0 Y-1995	la, Byrne, Bai l, Stewart & O Farm Road	n US/0844; , Robert (, Jeffrey (Nucleic) Controllic (Specific) 76
1ec	08/245 1994 1ON: E M. 24,02 ER: 2 MATION	DOS 5.1 8/8/08	Byrne Stewari	US/084 Robert Jeffre ucleic ontrol
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e	,356 5 7101	4 6	P 20 101	J., A., Acid Acid ling 1
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	5,356 25 271010-360)9A		. US/08442809A Robert J., Jeffrey A. Nucleic Acid Sequentrolling Lung Specific Gene Exp 76
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US-09-583-610D-17
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US-09-396-196G-81113
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Best Local S
Matches 15
                                                                Query Match
Best Local S
Matches 14
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SEQ ID NO 81113
LENGTH: 25
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Best Local Similarity
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SEQ ID NO 17
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                                                                                                                                                                                                                                                                                                 APPLICANT: Gianturco, S.H.
APPLICANT: Bradley, W.A.
APPLICANT: Bradley, W.A.
TITLE OF INVENTION: DNA Encoding Human apoB48R: A Monocyte-Macrophage
TITLE OF INVENTION: Apolipoprotein B48 Receptor Gene and Protein
FILE REFERENCE: D5880CIP
CURRENT APPLICATION NUMBER: US/09/583,610D
CURRENT FILING DATE: 2000-05-31
PRIOR APPLICATION NUMBER: US 09/130,242
PRIOR FILING DATE: 1998-08-06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Affymetrix, Inc.
TITLE OF INVENTION: Methods of Genetic Analysis
FILE REFERENCE: 3101.1
CURRENT APPLICATION NUMBER: US/09/396,196G
CURRENT FILING DATE: 1999-09-15
PRIOR PILING DATE: 1998-09-17
RIOR PILING DATE: 1998-09-17
NUMBER OF SEQ ID NOS: 127806
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Michael Mittmann APPLICANT: David Mack
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: DNA
                                                                                                                                              OTHER INFORMATION: Antisense primers were based on OTHER INFORMATION: sequence of the THP-1 (73 clone OTHER INFORMATION: of ApoB48R cDNA)
                                                                                                                                                                                                    TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                   PEATURE:
                                                                                                                                                                                                                                                   ENGTH: 17
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                                                            h 71.1%; Score 12.8; DB 3;
Similarity 87.5%; Pred. No. 2.1e+03;
14; Cohservative 0; Mismatches 2;
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 CCAGCCTCTTCTGGAT 16
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ilarity 93.8%;
Conservative
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Pred. No. 3.4e+02;
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Pred. No. 1.1e+03;
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US-09-396-196G-56205/c
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PRIOR FILING DATE: 1998-09-17
NUMBER OF SEQ ID NOS: 127806
SOPTWARE: FRAESEQ for Windows Version 4.0
SEQ ID NO 56205
LENGTH: 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
APPLICANT: Michael Mittmann
APPLICANT: David Mack
APPLICANT: David Lockhart
APPLICANT: Affymetrix, Inc.
TITLE OF INVENTION: Methods of Genetic Analysis
FILE REFERENCE: 3101.1
CURRENT APPLICATION NUMBER: US/09/396,196G
CURRENT FILING DATE: 1999-09-15
CURRENT FILING DATE: 1999-09-15
GENERAL INFORMATION:
APPLICANT: PROCKOP
APPLICANT: SPOTILA
APPLICANT: DELTAS,
APPLICANT: SEREDA,
APPLICANT: LARSON,
                                                                                                                 Sequence 305, Application US/08943731 Patent No. 6265157
                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQ ID NOS: 6849
SEQ ID NO 2493
LENGTH: 20
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APPLICANT: Griffais,
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Patent No. 6559294
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                                                                                                                                                                                                                                                                                                                                                                                                             ORGANISM: Chlamydia pneumoniae
                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: DNA
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PROCKOP, DARWIN J.
SPOTILA, LORETTA D.
DELTAS, CONSTANTINOS D.
SEREDA, LARISA
LARISA
LARSON, ANDREA W.
                                                                                                                                                                                                                                                                                                              Conservative
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                                                                                                                                                                                                                                                                                                                                 68.9%; Score 12.4; DB 3; 92.9%; Pred. No. 3.5e+03;
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APPLICANT

PACK, MICHAEL

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Sequence 24325, Application US/09396196G
Patent No. 6821724
GENERAL INFORMATION:
APPLICANT: Michael Mittmann
APPLICANT: David Mack
APPLICANT: David Lockhart
APPLICANT: NUMBER: US/09/396,1966
TITLE OF INVENTION: Methods of Genetic Analysis
FILE REFERENCE: 3101.1
CURRENT APPLICATION NUMBER: US/09/396,196G
CURRENT FILING DATE: 199-09-15
PRIOR APPLICATION NUMBER: 60/100,678
PRIOR FILING DATE: 1998-09-17
NUMBER OF SEQ ID NOS: 127806
SOFTWARE: FastSEQ for Windows Version 4.0
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US-09-396-196G-24325
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TELEPHONE: 215-965-1284
TELEPHONE: 25-567-2991
TELEX: 831<sup>1</sup>494
INFORMATION FOR SEQ ID NO: 305:
SEQUENCE CHARACTERISTICS:
LENGTH: 25 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOPTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/943,731
PILING DATE: 03-OCT-1997
CLASSIPICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/212,322
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/212,322
PRIOR DATE: 14-MAR-1994
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APPLICATION NUMBER: US 07/803,628
FILING DATE: 03-DEC-1991
ATTORNEY/AGENT INFORMATION:
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TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR DETECTING
TITLE OF INVENTION: ALTERED TYPE I OR TYPE IX COLLAGEN GENE SEQUENCES
NUMBER OF SEQUENCES: 666
CORRESPONDENCE ADDRESS:
ADDRESSEE: PANITCH SCHWARZE JACOBS & NADEL, P.C.
STREET: ONE COMMERCE SQUARE, 2005 MARKET STREET, 22ND
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MEDIUM TYPE: Floppy disk
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: DOYLE LEARY Ph.D., REGISTRATION NUMBER: 36, REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Match 68.9%;
Local Similarity 92.9%;
tes 13; Conservative
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TOPOLOGY: linear
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EARLY, JAMES
KORKKO, JARMO
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Pred. No. 3.6e+03;
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RESULT 9
US-07-722-798A-77
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; LENGTH: 25
; TYPE: DNA
; ORGANISM: mus musculus
US-09-396-196G-45539
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION NUMBER: 60/100,678
PRIOR FILING DATE: 1998-09-17
NUMBER OF SEQ ID NOS: 127806
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 45539
LENGTH: 25
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Best Local Similarity
Matches 13; Conserva
                                                                                                                                                                                                                                                                                                                                                          Patent No.
                                                                                                                                                                                                                                                                                                                                                                         Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Michael Mittmann
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Patent No. 682172
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: David Mack
APPLICANT: David Lockhart
APPLICANT: Affymetrix, Inc.
TITLE OF INVENTION: Methods of Genetic Analysis
FILE REFERENCE: 3101.1
CURRENT APPLICATION NUMBER: US/09/396,196G
CURRENT FILING DATE: 1999-09-15
                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
APPLICANT: Larry
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LENGTH: 25
TYPE: DNA
ORGANISM: Mus musculus
                                                                                                                                                                                              APPLICANT:
APPLICANT:
APPLICANT:
                                                                                                                                         APPLICANT: Natalie A. SOLOMON
TITLE OF INVENTION: AMPLIFICATION OF TARGET NUCLEIC
TITLE OF INVENTION: ACIDS USING GAP FILLING LIGASE CHAIN REACTION
NUMBER OF SEQUENCES: 124
                                                                                                                                                                                                                                             APPLICANT:
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy diskette
                                                                                                                               CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                             APPLICANT:
                                                                                                                                                                                                                                                                                              APPLICANT:
                                                                                                                                                                                                                                                                                                               APPLICANT:
                                                                 ADDRESSEE: Abbott Laboratories
STREET: One Abbott Park Road
CITY: Abbott Park
STATE: Illinois
                                                  COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                           Application US/07722798A
                                                                                                                                                                                                            Hsiang-Yun Hu
Jon David Kratochvil
Thomas G. Laffler
Ronald L. Marshall
Laurie A. Rinehardt
                                                                                                                                                                                                                                                                                            John J. Carrino
Bruce J. Dille
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                          Larry G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Application US/09396196G
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92.9%;
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Pred. No. 3.6e+03;
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Pred. No. 3.
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0,

COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOUTWARE: WOO'dperfect 5.0
CURRENT APPLICATION DATA:

APPLICATION NUMBER: FILING DATE: 199106 CLASSIFICATION: 435

19910628

US/07/722,798A

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Sequence 2002, Application US/09396196G

Patent NO. 6821724

GENERAL INFORMATION:
APPLICANT: Michael Mittmann
APPLICANT: David Mack
APPLICANT: David Lockhart
APPLICANT: David Lockhart
APPLICANT: Affymetrix, Inc.
TITLE OF INVENTION: Methods of Genetic Analysis
FILE REFERENCE: 3101.1
CURRENT APPLICATION NUMBER: US/09/396,196G
CURRENT APPLICATION NUMBER: 00/100,678
PRIOR APPLICATION NUMBER: 60/100,678
PRIOR APPLICATION NUMBER: 60/100,678
PRIOR APPLICATION NUMBER: 60/100,678
PRIOR FILING DATE: 1998-09-17
NUMBER OF SEQ ID NOS: 127806
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 20092
LENGTH: 25
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US-09-396-196G-20092/c
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                                                                                                                          RESULT 11
US-09-396-196G-20103/c
                                                                                                                                                                                                                                                                                                                                              US-09-396-196G-20092
GENERAL INFORMATION:
APPLICANT: Michael Mittmann
APPLICANT: David Mack
APPLICANT: David Lockhart
APPLICANT: Affymetrix, Inc
                                                                                      Sequence 20103, Application US/09396196G Patent No. 6821724
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                     ORGANISM: Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELECOMMUNICATION INFORMATION:
TELEPHONE: 708-937-4884
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MOLECULE TYPE: Other nucleic acid (synthetic DNA)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STRANDEDNESS: BILL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME: Thomas D. Brainard
REGISTRATION NUMBER: 32,459
REFERENCE/DOCKET NUMBER: 4773.US.P2
                                                                                                                                                                                                                                                                                             Local
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David Lockhart
David Lockhart
Affymetrix, Inc.
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82.4%;
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82.4%;
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Pred. No. 4.6e+03;
0; Mismatches 3;
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Pred. No. 4.6e+03;
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RESULT 13

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RESULT 12
US-08-480-473B-28/c
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; ORGANISM: Mus musculus
US-09-396-196G-20103
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                                                                                                                              US-08-480-473B-28
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LENGTH: 25
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CURRENT FILING DATE: 1999-09-15
FRIOR APPLICATION NUMBER: 60/100,678
PRIOR FILING DATE: 1998-09-17
NUMBER OF SEQ ID NOS: 127806
SOFTWARE: FASTSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
APPLICANT: Semenza, Gregg L.
TITLE OF INVENTION: HYPOXIA INDUCIBLE FACTOR-1 AND METHOD OF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILE REFERENCE: 3101.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: Methods of Genetic Analysis
                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US/08/480
FILING DATE: 06-JUN-1995
CLASSIFICATION: 514
ATTORNEY/AGENT INFORMATION:
NAME: Halle, Liea A.
REGISTRATION NUMBER: 38,347
                                                                                                                                                                                                                                           TELEPHONE: 619/678-50
TELEPAX: 619/678-5099
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                            STRANDEDNESS: 811
TOPOLOGY: linear
MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                                           REFERENCE/DOCKET NUMBER: 07
TELECOMMUNICATION INFORMATION:
TELEPHONE: 619/678-5070
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQUENCES:
                                                                                                                                                                                                                           SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STREET: 42
CITY: La J
STATE: CA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Match 67.8%;
Local Similarity 82.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADDRESSEB:
                                                                               Local
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                                1 CCCCCCTCTTCTGGATA 17
                                                                14;
                                                                               Similarity
                                                                                                                                                                                           1: 27 base pairs nucleic acid
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                                                              Conservative
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                                                                                                                                                                            single
                                                                               67.8%;
82.4%;
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Pred. No. 4.6e+03;
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                                                                               Score 12.2; DB 2;
Pred. No. 4.7e+03;
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US-08-915-213-28/c
                                                                                                                                                                                                      Patent No. 6020462
GENERAL INFORMATION:
                                                                                                                                                                                                                            Sequence 28, Application US/08915213 Patent No. 6020462
ZIP: 92037
COMPUTER READABLE FORM:
                                                                                                                                  APPLICANT: Semenza, Gregg L.
TITLE OF INVENTION: HYPOXIA INDUCIBLE FACTOR-1 AND METHOD OF USE
NUMBER OF SEQUENCES: 64
CORRESPONDENCE ADDRESS:
                                                  STREET: 4225 Executive Square, Suite 1400 STREE: La Jolla STATE: CA.
                                              COUNTRY: USA
```

US-08-915-213-28 Matches Query Match SEQUENCE CHARACTERISTICS:
LENGTH: 27 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear TELEPAX: 619/678-5099 INFORMATION FOR SEQ ID NO: NAME: Haile, Lisa A.
REGISTRATION NUMBER: 38,347
REFERENCE/DOCKET NUMBER: 07265/053001
TELECOMMUNICATION INFORMATION:
TELEPHONE: 619/678-5070 MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION UNMERR: US/08/915,213
FILING DATE: 20-AUG-1997 TOPOLOGY: 11 ATTORNEY/AGENT INFORMATION: CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION UNMBER: US 08/480,473
PILING DATE: 06-JUN-1995 Local Similarity 82.4 27 1 CCCCCCTCTCTGGATA 17 CCCCCCTTTTCAGAATA 11 67.8**%**; 82.4**%**; 28: 0 Score 12.2; DB 3; Pred. No. 4.7e+03; Mismatches DB 3; Length 27; Indels 0

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US-09-235-217-28/c Sequence 28, Application US/09235217 Patent No. 6222018 GENERAL INFORMATION: SOPTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA: COMPUTER READABLE FORM: MEDIUM TYPE: Floppy CORRESPONDENCE ADDRESS NUMBER OF SEQUENCES: APPLICANT: Semenza, TITLE OF INVENTION: MEDIUM TYPE: Ploppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS COUNTRY: U ZIP: 92037 STATE: STREET: ADDRESSEE: La Jolla S: CA B: Fish & Richardson P.C 4225 Executive Square, St USA , Gregg L.
HYPOXIA INDUCIBLE FACTOR-1 AND METHOD OF USE Square, Suite 1400

APPLICATION NUMBER:

US/09/235,217

Query Match Matches

Local Similarity les 14; Conserv

Conservative

67.8%;

Score 12.2; Pred. No. 4.

Mismatches

4.7e+03; 3;

Indels

0

Gaps

0

Length 27;

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                        S
                                                                                                            US-09-235-217-28
                                                     Matches
                                                                  Query Match
Best Local S
                                                                                                                                                                                        TELEFAX: 619/678-5099
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                              ATTORNEY/AGENT INFORMATION:
NAME: Haile, Lisa A.
REGISTRATION NUMBER: 38,347
REFERENCE/DOCKET NUMBER: 07:
TELECOMMUNICATION INFORMATION:
                                                                                                                          MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION DATA:
                                                                                                                                      LENGTH: 27 base pairs TYPE: nucleic acid STRANDEDNESS: single TOPOLOGY: linear
                                                                   Local Similarity
                                                                                                                                                                                                                                    TELEPHONE:
                                                                                                                                                                                                                                                                                                                      FILING DATE:
                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                CLASSIFICATION:
27
                                                  14;
                        1 CCCCCCTCTTCTGGATA 17
CCCCCTTTCAGAATA 11
                                                     Conservative
                                                                                                                                                                                                                                     619/678-5070
                                                                                                                          DNA
                                                                67.8%;
82.4%;
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                                                   0,
                                                                                                                                                                                                                                                               07265/053001
                                                               Score 12.2; DB 3;
Pred. No. 4.7e+03;
                                                   Mismatches
                                                                           Length 27;
                                                   Indels
                                                  o,
                                                 Gaps
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RESULT 15 PCT-US96-10251-28/c

0

PCT-US96-10251-28 Sequence 28, Application PC/TUS9610251 GENERAL INFORMATION: ATTORNEY/AGENT INFORMATION:
NAME: Haile, Lisa A.
REGISTRATION UNMER: 38,347
REFERENCE/DOCKET NUMBER: 0726
TELEPOMUNICATION INFORMATION:
TELEPIONE: 619/678-5070
TELEPAX: 619/678-5099 TELEFAX: 619/678-5099 INFORMATION FOR SEQ ID NO: STRANDEDNESS:
TOPOLOGY: line
MOLECULE TYPE: 1 SOPTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US96/10251
FILING DATE: 06-JUN-1996
CLASSIFICATION: SEQUENCE CHARACTERISTICS: LENGTH: 27 base pairs TYPE: nucleic acid COMPUTER READABLE FORM:
MEDIUM TYPE: Ploppy of COMPUTER: IBM PC comp APPLICANT: The Johns Hopkins University School of Medicine TITLE OF INVENTION: HYPOXIA INDUCIBLE FACTOR-1 AND METHOD NUMBER OF SEQUENCES: CORRESPONDENCE ADDRESS STATE: C MEDIUM TYPE: Ploppy disk COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS CITY: La Jolla STATE: CA STREET: ADDRESSEE: 92037 4225 Executive Square, Suite 1400 USA Fish & Richardson P.C. single 07265/053WO1 OF USE

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Result
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Maximum Match 100%
Listing first 45 summaries
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Match Length DB
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US-10-659-980A-5
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gb_est6:*
gb_est7:*
gb_gss1:*
gb_gss2:*
gb_gss3:*
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gb_est2:*
gb_est3:*
gb_htc:*
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      AZ592213
AI915575
AI649190
CX001953
AZ584820
AZ465690
AZ660176
CX007773
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AZ487733
AI039313
AZ875769
AZ627859
AZ585557
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     AZ785027 2M0028H03
BQ589768 E012680-0
AZ487733 1M0317A10
AI039313 0x36b04.8
AZ875769 2M0190A02
AZ627575 1M0390M18
D18736 MUSGS01798
AZ558557 1M0390M18
D18736 MUSGS01798
AZ592213 1M0402P23
AI915575 tr40g99.x
AI649190 uk22d12.x
CX001953 iv44a06.b
AZ584820 1M0389A14
AZ465690 1M0278910
AZ660176 1M0389A12
AZ660176 1M0339A12
AZ66017773 io30009.b
                                                                                                                                                    AL453426 T. brucei
AZ817329 2M0086D22
R54679 y974e12.rl
AZ489682 1M0322G16
CZ443094 IBB8D03.f
                                                                                                                                                                                                           Description
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ALIGNMENTS

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REFERENCE
AUTHORS
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VERSION
KEYWORDS
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TA25G06P/c
LOCUS
DEFINITION
                                                                                            ORIGIN
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                                                                                                                                                                                                                                                      FEATURES
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Query Match 67.8%;
Best Local Similarity 82.4%;
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE
JOURNAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ORGANISM
                                                                                                                                                                                                                                source
                                                                                                                                                                                                                                          Constructed at the Institute for Genomic Research (TIGR),
Rockville, MD. Genomic DNA isolated from a cloned population of
Trypanosoma brucei (TREU927/4 GUTat 10.1) was mechanically sheared
to give a tight size distribution (
4 kb). The v + i method used for the library construction is
described in detail in Smith, H. and Venter, J.C. (Making small
insert libraries for whole genome shotgun sequencing projects. In
Genome Sequencing: A practical Approach, eds. M. Vaudin and B.
Barrell, Oxford University Press, 1999).
Email: nelsayed@tigr.org
Details of T. brucei sequencing at the Sanger Centre are available
at http://www.sanger.ac.uk/Projects/T_brucei/.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Direct Submission
Submitted (10-DEC-2000) Trypanosoma brucei genome sequencing project, Sanger Centre, The Wellcome Trust Genome Campus, Hinxton, Cambridge CB10 1SA, E-mail: barrell@sanger.ac.uk and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Hall, N., Bowman, S., Lennard, N.J., Doggett, J., Atkin, R., Chillingworth, C., Ormond, D., Harris, B., El-Sayed, N., Hou, L., Melville, S.E., Rajandream, M.A. and Barrell, B.G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Trypanosoma brucei
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AL453426.1 GI:11849213
GSS.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Eukaryota; Euglenozoa; Kinetoplastida; Trypanosomatidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (bases 1 to 26)
                                                                                                            /organism="Trypanosoma brucei"
/mol type="genomic DNA"
/strāin="TREU927"
/db xref="taxon:5691"
/clone="25g06"
Score 12.2; DB 11;
Pred. No. 2.5e+05;
0; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          26 bp D
DNA clone
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1e 25g06,
                                          Length 26;
  Indels
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Gaps
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ORIGIN
Query Match
Best Local Similarity
                                                                                                                                         was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was hint end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoreeis. Vector DNA was prepared from a derivative of pWD42 (gi | 4732114 | gb | AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
                  65.6%;
                  Score 11.8; DB 9;
Pred. No. 3.8e+05;
                                                   Length 22;
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2 CCCCCTCTTCTGGAT 16

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Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R. Mouse whole genome scaffolding with paired end reads from 10kb Nouse whole genome scaffolding with paired end reads
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia;
Sciurognathi; Muroidea; Muridae; Murinae; Mus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AZ817329 22 bp DNA linear GSS 20
2M0086D22R Mouse 10kb plasmid UUGCIM library Mus musculus
clone UUGC2M0086D22 R, genomic survey sequence.
                                                                                                                                                                                                                                                                                                                                                               /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"/clone lib="Mouse 10kb plasmid UUGCIM library"/note="Vector: PWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackson Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              S. 2030 E., SLC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GSS 20-FEB-2001
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R54679/c
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AUTHORS
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               Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Email: est@watson.wustl.edu
Insert Size: 718
High quality sequence starts: 1 High quality sequence stops: 1
Source: IMAGE Consortium, LLNL This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further information. Trace considered overall poor quality Possible reversed clone: similarity on wrong strand
Insert Length: 718
Std Error: 0.00
Seg primer: M13RP1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 (bases 1 to 28)
Hillier,L., Clark,N., Dubuque,T., Elliston,K., Hawkins,M.,
Holman,M., Hultman,M., Kucaba,T., Le,M., Lennon,G., Marra,M.,
Parsons,J., Rifkin,L., Rohlfing,T., Soares,M., Tan,F.,
Trevaskis,B., Waterston,R., Williamson,A., Wohldmann,P. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 R54679

Yj74e12.r1 Soares breast 2NbHBst Homo sapiens cDNA clone
Yj74e12.r1 Soares breast 2NbHBst Homo sapiens cDNA clone
IMAGE:154510 5' similar to SP:A47291 A47291 PUTATIVE ZINC FINGER
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Washington University School of Medicine 4444 Forest Park Parkway, Box 8501, St. I Tel: 314 286 1800
Fax: 314 286 1810
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Unpublished (1995)
Contact: Wilson RK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The WashU-Merck EST Project
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens (human)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PROTEIN - ;, mRNA sequence.
                        Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CCCCCTTTTCTGAA 22
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Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            quality sequence stop: 1.
Location/Qualifiers
                                                                                                       /organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="GDB:566948"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                       /sex="Female"
                                                                                                                                                                                                                                                                                                                                                                                                                /clone="IMAGE:154510"
                    65.6%;
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  0
  Score 11.8; DB 8;
Pred. No. 3.9e+05;
0; Mismatches 2;
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                                          Length 28;
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    Gaps
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JOURNAL COMMENT

plasmid inserts
Unpublished (2000)
Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah

308,

Biomedical Polymers Research Bldg., 20

USA

TITLE

FEATURES

High quality sequence stop: 22.
Location/Qualifiers

/mol_type="genomic DNA"
/strain="C57BL/6J"
/db_xref="taxon:10090" organism="Mus musculus"

/sex="Male"

/clone="UUGC2M0086D22"

ass: plasmid ends

Tel: 801 585 5606

Fax: 801 585 7177

Email: ddunn@genetics.utah.edu

Insert Length: 10000 Std Error: 0.00

Plate: 0086 row: D column: 22

Seq primer: CACACAGGAAACAGCTATGACC

Bource

REFERENCE AUTHORS

(bases 1 to 22)

ACCESSION VERSION KEYWORDS

AZ817329.1 GI:12987333 Mus musculus (house mouse)

ORGANISM

Mus musculus

RESULT 2 AZ817329

DEFINITION

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25

CCCCCCTCTTCATGAAA 9 CCCCCCTCTTCTGGATA 17

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18

CCCCCCACTTCTAGA

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                                                   Query Match
Best Local Similarity
Matches 13; Conserv
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1 CCCCCCTCTTCTGGA 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Brror: 0.00
Plate: 0322 row: G column: 16
Seq primer: CGTTGTAAAACGACGGCCAGT
Class: Dlasmid ende
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weise, R. Mouse whole genome scaffolding with paired end reads from 10kb Mouse whole genome scaffolding with paired end reads
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              University of Utah Genome Center University of Utah Rm. 308, Biomedical Polymers Res
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        plasmid inserts
Unpublished (2000)
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28 bp DNA line 1M0322G16F Mouse 10kb plasmid UUGC1M library Muclione UUGC1M0322G16 F, genomic survey sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    High quality sequence stop: 28.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Contact: Robert B. Weiss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sciurognathi; Muroidea; Muridae; Murinae; Mus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Eukaryota;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (bases 1 to 28)
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                                                   Conservative
                                                                                                                                                                  (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWH2 (gi |4732114 |gb|AF129072.1), a copy-number inducible derivative of plasmid Rl. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent B. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /lab_host="B. Coli strain XL10-Gold, T1-resistant, F-"/Clone_11b="Mouse 10kb plasmid UUGCIM library"/note="Vector: pW042nv; Purified genomic DNA from M.musculus C57BL/6J (male) was obtained from the Jackson
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/strain="C57BL/6J"
/db_xref="taxon:10090"
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                                                                           65.6%;
86.7%;
                                                0,
                                                                   Score 11.8; DB 9;
Pred. No. 3.9e+05;
                                                   Mismatches
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                                                                                              Length 28;
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PUBMED
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1 (bases 1 to 28)

1 (bases 1 to 28)

Lewinski, M.K., Bisgrove, D., Shinn, P., Chen, H., Hoffmann, C.,

Lewinski, M.K., Bisgrove, D., Shinn, P., Chen, H., Hoffmann, F.D.

Hannenhalli, S., Verdin, E., Berry, C.C., Ecker, J.R. and Bushman, F.D.

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Hannenhalli, S., Werth, E., Berry, C.C., Ecker, J.R. and Bushman, F.D.

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Fax: 215 573 4856
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CZ443094.1 GI:62379366
GSS.
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University of Pennsylvania School of Medicine
402C Johnson Pavilion, 3610 Hamilton Walk, Philadelphia, PA
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Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini;
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that are associated with repression of an integrated human immunodeficiency virus type 1 (HIV-1) promoter. HIV-1 integrated not results in the formation of an integrated human immunodeficiency virus type 1 (HIV-1) promoter. HIV-1 integration results in the formation of precise and homogeneous junctions between viral and hose DNA, but integration takes place at many locations. Thus, the variation in HIV-1 gene expression at different integration sites reports the activity of regulatory sequences at nearby chromosomal positions. Negative regulation of HIV transcription is of particular interest because of its association with maintaining HIV in a latent state in cells from infected patients. To identify chromosomal regulators of HIV transcription, we infected Jurkat T cells with an HIV-based vector transducing green fluorescent protein (GFP) and separated cells into populations containing well-expressed (GFP-positive) or poorly expressed (GFP-negative) proviruses. We then determined the chromosomal locations of the two classes by sequencing 971 junctions between viral and cellular DNA. Possible effects of endogenous cellular transcription were characterized by transcriptional profiling. Low-level GFP expression correlated with integration in (1) gene deserts, (11) centromeric heterochromatin, and (111) very highly expressed cellular genes. These data provide a genome-wide picture of chromosomal features that repress transcription and suggest models for transcriptional latency in cells from HIV-infected patients."
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/db_xref="taxon:9606"
/clone="IBB8D03.fwd"
/cell_line="Jurkat"
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JOURNAL COMMENT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Tel: 801 585 5606

Pax: 801 585 7177

Email: Bdunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00

Plate: 0028 row: H column: 03

Seg primer: CACACAGGAAACAGCTATGACC

Class: plasmid ends
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, B., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Unpublished (2000)
Contact: Robert B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mouse whole genome scaffolding with paired end reads from 10kb plasmid inserts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eut
Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia;
Sciurognathi; Muroidea; Muridae; Murinae; Mus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AZ785027.1
GSS.
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2M0028H03R Mouse 10kb plasmid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              University of Utah Genome Center
University of Utah
Rm. 308, Biomedical Polymers Research Bldg., 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Class: plasmid ends
High quality sequence stop: 23.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CCCCTCTTCTGGATA 17
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                                                             Laboratory Mouse DNA Resource (http://www.jax.org/resources/documents/dnaxes/). The DNA (http://www.jax.org/resources/documents/dnaxes/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gi|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptore ONA, and transformed into
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ÷
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                                                                                                                                                                                                                                                                                                                                                                                                                                           /lab_host="B. Coli strain XL10-Gold, Tl-resistant, F-" /clone lib="Mouse 10kb plasmid UUGCIM library" /notee="Vector: pWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /mol_type="genomic DNA"
/strain="C57BL/6J"
/db_xref="taxon:10090"
                    chemically-competent E. coli XI10-Gold (Stratagene) cells and selected for ampicillin resistance."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /gex="Male"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /clone="UUGC2M0028H03"
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86.7%;
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Pred. No. 3.9e+05;
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d UUGC1M library Mus musculus genomic
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PUBMED
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Best Local
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                                                                                                  Best Local
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B012680-024-020-D03-SP6 MPIZ-ADIS-024-storage root
CDNA Clone 024-020-D03 5-PRIME, mRNA sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Insert Length: 13 Std Error: 0.00
Plate: 20 row: D column: 03
Seq primer: SP6; CATACGATTTAGGTGACACTATAG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADIS DNA core facility at MPIZ
Max-Planck-Institute for Plant Breeding Research
Carl-von-Linne Weg 10, 50829 Koeln, Germany
Fax: 00492215062851
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Construction of a 'unigene' cDNA clone set by oligonuclectide fingerprinting allows access to 25 000 potential sugar beet genes plant J. 32 (5), 845-857 (2002)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Herwig,R., Schulz,B., Weisshaar,B., Hennig,S., Steinfath,M., Drungowski,M., Stahl,D., Wruck,W., Menze,A., O'Brien,J., Leb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Bukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicotyledons; Caryophyllales; Amaranthaceae; Beta.

1 (bases 1 to 13)
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                                        4 CCCTCTTCTGGAT 16
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                                                                                                  Similarity
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CCCTCTTCTTGAT 13
                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    weisshaa@mpiz-koeln.mpg.de
                                                                                                                                                                                                                                                                                                        /clone_lib="MPIZ-ADIS-024-storage root"
/note="Vector: pCMVSPORT6; Site 1: SalI; Site 2: NotI;
/note="Vector: pCMVSPORT6; Site 1: SalI; Site 2: NotI;
cDNA library from sugar beet, llbrary provided by KWS
Kleinwanzlebener Saatzucht AG Einbeck, Germany, contact
b.schulz@kws.de; cloning sites SalI-NotI, primer sites
                                                                                                                                                                                                       Sequencing granted in the context of the GABI-Beet project, local PI: Dr. Katharina Schneider, coordinator: Prof. Christian Jung; Sequence submission managed by
                                                                                                                                                                                                                                                  SP6-SalI-CCACGCGTCCG-Sprime-cDNA-polyA-CC-NotI-T7; Note: Sequencing granted in the context of the GABI-Beet
                                                                                                                                                                                                                                                                                         orientation:
                                                                                                                                                                                                                                                                                                                                                                                                             /db xref="taxon:161934"
/clone="024-020-D03"
/tissue_type="storage root"
/lab_host="EMDH10B"
                                                                                                                                                                                    RZPD/GABI-Primary database: http://gabi.rzpd.de"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          cultivar="KWS2320 (double haploid, monogerm breeding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              organism="Beta vulgaris"
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                                                                                                  63.3%;
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Pred. No. 4.8e+05;
                                                                             Score 11.4; DB 5;
Pred. No. 5.7e+05;
D; Mismatches 1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Brror: |
Plate: 0317 row: A column: 10
Seq primer: CACACAGGAAACAGCTATGACC
Class: Diasmid ends
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           University of Utah Genome Center
University of Utah
Rm. 308, Biomedical Polymers Research Bldg., 20 S. 2030 B., SLC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                plasmid inserts
Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia;
Sciurognathi; Muroidea; Muridae; Murinae; Mus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   clone UUGC1M0317A10
AZ487733
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AZ487733 26 bp DNA linear GSS 05-OCT-2001M0317A10R Mouse 10kb plasmid UUGCIM library Mus musculus genomic clone UUGCIM0317A10 R, genomic survey sequence.
                                                                                                                                                                                                           Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Contact: Robert B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mus musculus (house mouse)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AZ487733.1 GI:10655751
                                                CCCTCTTCTGGAT 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (bases 1 to 26)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  quality sequence stop: 26.
Location/Qualifiers
                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                        (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pW142 (gi |4732114 |gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent B. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /lab_host="B. Coli strain XL10-Gold, T1-resistant, F-"/clone_lib="Mouse_10kb_plasmid_UUGCIM_library"/note="Vector: PWD42nv, Purified_genomic_DNA_from_M.musculus_C57BL/6J_(male)_was_obtained_from_the_Jackson_musculus_C57BL/6J_(male)_was_obtained_from_the_Jackson_musculus_C57BL/6J_(male)_was_obtained_from_the_Jackson_musculus_C57BL/6J_(male)_was_obtained_from_the_Jackson_musculus_C57BL/6J_(male)_was_obtained_from_the_Jackson_musculus_C57BL/6J_(male)_was_obtained_from_the_Jackson_musculus_C57BL/6J_(male)_was_obtained_from_the_Jackson_musculus_C57BL/6J_(male)_was_obtained_from_the_Jackson_musculus_C57BL/6J_(male)_was_obtained_from_the_Jackson_musculus_C57BL/6J_(male)_was_obtained_from_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_the_Jackson_
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/clone="UUGC1M0317A10"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /mol_type="genomic DNA"
/strain="C57BL/6J"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              organism="Mus musculus"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   sex="Male"
                                                                                                                                                                                                           63.3%;
92.3%;
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                                                                                                                                                              0
                                                                                                                                                                                                  Score 11.4; DB 9;
Pred. No. 6.1e+05;
                                                                                                                                                      Mismatches
                                                                                                                                                                                                                                                  DB 9;
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8
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REFERENCE
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AZ875769
LOCUS
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VERSION
KEYWORDS
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TITLE
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AI039313
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VERSION
KEYWORDS
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Best Local !
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  13;
Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia;
Sciurognathi; Muroidea; Muridae; Murinae; Mus.
1 (Dases 1 to 19)
Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C.,
Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T.,
                                                                                                                                                                                                               19 bp DNA linear GSS 21 2M0190A02R Mouse 10kb plasmid UUGC1M library Mus musculus clone UUGC2M0190A02 R, genomic survey securence AZ875769
                                                                                                                                                                        GSS.
                                                                                                                               Mus musculus
                                                                                                                                                                                            AZ875769.1 GI:13086107
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Contact: Robert Strausberg, Ph.D.

Email: cgapbs-r@mail.nih.gov

This clone is available royalty-free through LLNL; contact the

IMAGE Consortium (info@image.llnl.gov) for further information.

Insert Length: 954 Std Error: 0.00

Seq primer: -40ml3 fwd. ET from Amersham
                                                                                                                                                 Mus musculus (house mouse)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AIO39313 22 bp mRNA linear SGT 24.SEP-199 0x36b04.81 Soares total fetus Nb2HF8 9w Homo sapiens cDNA clone IMAGE:1658383 3' Similar to SW:TSP3 HUMAN P49746 THROMSOSPONDIN 3 PRECURSOR. ; contains Alu repetitive element; mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Hominidae; Homo.
1 (bases 1 to 22)
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                                                                                                                                                                                                                                                                                                                                                                               CCCCCCTCGCATGGAT 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Double-stranded cDNA was ligated to Eco RI adaptors (Pharmacia), digested with Not I and cloned into the Not I and Eco RI sites of the modified pT773 vector. Library went through one round of normalization, and was constructed by Bento Soares and M. Fatima Bonaldo. "
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /clone="IMAGE:1658383"
/dev_stage="8-9 weeks"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          62.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                             0,
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Pred. No. 7.6e+05;
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COMMENT
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AZ627859
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Insert Length: 10000 Std Error: (
Plate: 0190 row: A column: 02
Seq primer: CACACAGGAAACAGCTATGACC
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Fax: 801 585 7177
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University of Utah Genome Center
University of Utah
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Niederhausern,A. and Wright,D.,Weiss,R.
Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               High quality sequence stop: 19.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Unpublished (2000)
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                                                                                                                                                                                                                                                                                                            AZ627859 20 bp DNA linear GSS 13-DEC-200 1M0476E04F Mouse 10kb plasmid UUGCIM library Mus musculus genomic Clone UUGCIM0476E04 F, genomic survey sequence.
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                                                       Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eut
Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia;
Sciurognathi; Murcidea; Muridae; Murinae; Mus.
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                                                                                                                                                                                                                                                       AZ627859.1
                                                                                                                                                                                                                                                                                        AZ627859
Dunn, D., Aoyagi, A., Barber, M.,
                                                                                                                                                            Mus musculus
                                                                                                                                                                                      Mus musculus (house mouse)
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                                (bases 1 to 20)
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//Bab host="B. Coli strain XL10-Gold, T1-resistant, F-"
//Clone lib="Mouse 10kb plasmid UUGCIM library"
//Clone lib="Mouse 10kb plasmid UUGCIM library"
//Clone lib="Mouse 10kb plasmid UUGCIM library"
//Clone lib-"Mouse DNA resource
Intel://www.jax.org/resources/documents/dnares/). The DNA
musculius C57BL/6J (male) was obtained from the Jackson
Laboratory Mouse DNA resource
(http://www.jax.org/resources/documents/dnares/). The DNA
was hydrodynamically sheared by repeated passage through a
0.05 inch orifice at constant velocity. The sheared DNA
was blunt end-repaired with T4 DNA polymerase and T4
polynucleotide kinase. Adaptor oligonucleotides were
ligated to the blunt ends in high molar excess. The
adaptored DNA was purified and size-selected for a 9.5 to
10.5 kb range using preparative agarose gel
10.5 kb range using preparative agaro
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/strain="C57BL/6J"
/db_xref="taxon:10000"
/clone="UUGC2M0190A02"
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/mol_type="genomic DNA"
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      Beacorn, T.,
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         Duval, B.,
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             Hamil, C.,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Seq primer: CGTTGTAAAACGACGGCCAGT
Class: plasmid ends
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Err
Plate: 0476 row: E column: 0
Mus musculus
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Eukaryota; Metheria; Euarchontoglires; Glires; Rodentia;
Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia;
Sciurognathi; Muroidea; Muridae; Murinae; Mus.
1 (bases 1 to 25)
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1M0390M18R Mouse 10kb plasmid UUGC1M library Mus musculus clone UUGC1M0390M18 R, genomic survey sequence.
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University of Utah
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801 585 7177
                                                                                                                                        musculus (house mouse)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Laboratory Mouse DNA Resources (documents/dnares/). The DNA (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pNPA2 (gi|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent B. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
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/strain="C57BL/6J"
/db_xref="taxon:1000"
/clone="UUGC1M0476E04"
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100.0%; Pred. No. 9.4e+C
tive 0; Mismatches
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RESULT 13
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    Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Sciurognathi; Muroidea; Muridae; Murinae; Mus.
                                                                                                                                                                                                        D18736 26
MUSGS01798 Mouse 3'-directed
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                                                                         Mus musculus
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D18736
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                                                                                              Mus musculus (house mouse)
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Insert Length: 10000 Std Brror:
Plate: 0390 row: M column: 18
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Location/Qualifiers
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Seq primer: CACACAGGAAACAGCTATGACC
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Fax: 801 585 7177
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/strain="C57BL/6J"
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Mus musculus cDNA clone md0836 3'
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                                                                                                                                                                                                                                                                  Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0402 row: P column: 23
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                11;
                                                                                                                                                                                                    High quality sequence stop: 23.
                                                                                                                                                                                                                        Class: plasmid ends
                                                                                                                                                                                                                                           Seq primer: CACACAGGAAACAGCTATGACC
                                                                                                                                                                                                                                                                                                                                  Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                        Ra.
                                                                                                                                                                                                                                                                                                                                                                                                                          University of Utah University of Utah
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Maroidea; Muridae; Murinae; Mus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GSS.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Contact: Robert
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    plasmid inserts
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Email: shoko@next.imcb.osaka-u.ac.jp.
Location/Qualifiers
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Unpublished (1995)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Contact: Shoko Kawamoto
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CTCTTCTGGAT 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (bases 1 to 23)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                 USA
                                                                                                                                                                                                                                                                                                                                                                                                 Biomedical Polymers Research Bldg.,
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clone="UUGC1M0402P23"
                                                                                                                                                                            location/Qualifiers
lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"
                             sex="Male"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            tissue_type="decidual tissue (day 6.5-8.5 of gestation)"
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Pred. No.
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Length Indels

26

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linear

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SLC, S

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was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pwp42 (gi|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
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Laboratory_Mouse_DNA_Resource_
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RESULT 15 AI915575/c 밁 w

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CCTCTTCTGGATAC 18 CCTCTTCTGAACAC 16

ORIGIN

SOURCE ORGANISM ACCESSION VERSION DEFINITION KEYWORDS Bequence. AI915575

AI915575

LT40g09.x1 NCI CGAP Ov23 Homo sapiens cDNA clone IMAGE:2220832 3'

similar to SW:P100 HUMAN P30808 DNA-BINDING P52/P100 COMPLEX, 100

BURNALL CONTRACT CON Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini; Homo sapiens AI915575.1 GI:5635430 sapiens (human)

REFERENCE AUTHORS TITLE (bases 1 to 25)

Hominidae; Homo

NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap. National Cancer Institute, Cancer Genome Anatomy Project (CGAP),

COMMENT

JOURNAL

Tumor Gene Index
Unpublished (1997)
Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Christopher Moskaluk, M.D., Ph.D., Michael
Emmert-Buck, M.D., Ph.D.

cDNA Library Preparation: Life Technologies, Inc.
cDNA Library Arrayed by: Greg Lennon, Ph.D.
DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-CGAP clone distribution information can be
Found through the I.M.A.G.E. Consortium/LLNL at: information can be

www-bio.llnl.gov/bbrp/image/image.html

Trace considered overall poor quality Seq primer: -40UP from Gibco High quality sequence step: 1. Location/Qualifiers

PEATURES

source

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/site_2: NotI; Cloned unidirectionally. Primer: Oligo dT.
Average insert size 1.35 kb. Tumor types include: mixed
Mullerian tumor, papillary serous, clear cell, spindle
cell. All are primary tumors, metastasis positive. Life
Technologies catalog #: 11534-013"

Query Match Best Local S Matches 12 12; Similarity Conservative 60.0%; 85.7%; 0 Score 10.8; DB 1; Pred. No. 1.2e+06; 0; Mismatches 2; 2 Length 25; Indels 0 Gaps

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Search completed: April 7, 2006, 10:30:59 Job time: 2907.78 secs

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Minimum
Maximum
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Maximum Match 100%
Listing first 45 summaries
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2: geneseqn2900
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8: geneseqn2003
9: geneseqn2003
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12: geneseqn2000
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14: geneseqn2000
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geneseqn2003ds:*
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geneseqn2004bs:*
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Pred. No. is th score greater t and is derived is the number of results predicted by chance to have a ater than or equal to the score of the result being printed, rived by analysis of the total score distribution.

SUMMARIES

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12.4	12.6	12.8	12.8	12.8	12.8	13.2	13.2	13.2	13.2	13.2	13.2	13.4	13.8	13.8	14.4	18	18	Score
68.9	70.0	71.1	71.1	71.1	71.1	73.3	73.3	73.3	73.3	73.3	73.3	74.4	76.7	7.0.7	80.0	100.0	100.0	Query Match Length
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The invention relates to 4 novel types of ammonia-oxidising bacteria (AOB) found in freshwater aquaria. The bacteria are able to oxidise ammonia to nitrite and are members of the ammonia-oxidising bacteria family of the beta subdivision of Proteobacteria. The 4 types of bacterican de distinguished on the basis of their 16S rRNA (ribosomal RNA) gene

bacteria

New bacteria capable of oxidizing ammonia to nitrite, for preventing or alleviating the accumulation of ammonia in fresh water aquaria, seawater aquaria and waste water.

WPI; 2002-075367/10.

Hovanec TA,

Burrell

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Claim 26; Page 53; 62pp; English.

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12.2 12.2 12.2 12.2	12.2 12.2	12.2 12.2 12.2	12.2	12.2	12.4	12.4
67.8 67.8 67.8 67.8	67.8 67.8 67.8	67.8 67.8		67.8 67.8	67.8	0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0
2 2 3 3 3 3 4 3 3 3 3 3 3 3 3 3 3 3 3 3	220 21	19 19	19	17 18	25 17	222222
10 14	14 14 13	12 14	1116	10 34	, N 0 0	13
AAQ36081 AAX03938 ABK24426 ADE40344 ADW38589	ADZ84877 ADZ84729 ADU42656	ADQ14296 ADY87306 ADY87733	ADF36480 ADL79197 ADL78948	ADZ34093 AAA14813 ADF36053	ACI19119 AAA17514 ABZ65019	AAX93192 ABX03745 ABX13323 ADN02302 ADN02302 ADR68924
Aaq36081 HSV detec Aax03938 Herpes Si Abk24426 TrkC agon Ade40344 Forward A Adw38589 Hot flash	Human Human Knock-	Adq14296 CAPN3/DYS Ady87306 VEGFR sir Ady87733 VEGFR sir	Adf36480 Human VEG Adl79197 Human HER Adl78948 Human HER	Adz3403 Human HER Adz34093 Human HER Aaa14813 PCR prime Adf36053 Human VEG		Aax93192 PCR prime Abx03745 Human REC Abx13323 Human NSD Adn02302 PCR prime Adr68924 Human NSD

ALIGNMENTS

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RESULT 1
ABA02420
                                                                                                                                      19-MAY-2000; 2000US-00573684.
                                                                                                                                                    17-MAY-2001; 2001WO-US016265
                                                                                                                                                                      29-NOV-2001.
                                                                                                                                                                                                                  16S rRNA gene; ribosomal RNA; aquarium; aquacultu waste water treatment; bioremediation; probe; ss.
                                                                                                                                                                                     WO200190312-A1.
                                                                                                                                                                                                    Nitrosomonadales.
                                                                                                                                                                                                                                   Type A; type A1; ammonia-oxidising bacterium; AOB; nitrite;
                                                                                                                                                                                                                                                Type A/Al ammonia-oxidising bacterium 16S rRNA gene probe.
                                                                                                                                                                                                                                                                  29-AUG-2003
04-MAR-2002
                                                                                                                                                                                                                                                                                         ABA02420;
                                                                                                                                                                                                                                                                                                         ABA02420 standard; DNA; 18
                                                                                                                        (AQUA-)
                                                                                                                        AQUARIA INC.
                                                                                                                                                                                                                                                                  (revised)
(first entry)
                                                                                                                                                                                                                          ribosomal RNA; aquarium; aquaculture;
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RESULT 2
ADM32708
ID ADM3
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AC ADM3
XX
AC ADM3
AC ADM
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           19-SEP-2002;
19-SEP-2002;
19-SEP-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                New composition comprising an isolated bacterial strain that ammonia to nitrite, useful for alleviating or preventing the of ammonia in aqueous environment.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO2004026772-AE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Nitrosospira-like 16S rDNA probe.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADM32708;
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                                                              This sequence represents a probe which was used in the detection of 16S TDNA sequence derived from an ammonia oxidising bacteria (AOB). The amplified sequence may be used in a composition which comprises an isolated bacterial strain that oxidizes ammonia to nitrite. The composition may be used for alleviating or preventing the accumulation of ammonia in a medium. The ammonia is reduced by at least 30% when compared with a level of ammonia that would exist in the absence of the bacterial strain. The composition is useful for alleviating or preventing the accumulation of ammonia in aqueous environment, e.g. a freshwater or
                                                                                                                                                                                                                                                                                                                                                                                                     Example 6; Page 34; 98pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2004-304936/28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Hovanec
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   10-SEP-2003;
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aqueous environment; freshwater; seawater;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        17-JUN-2004
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2002US-0386218P.
2002US-0386219P.
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      S-G-Nsspa-0149-a-A-18.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        aquarium;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ammonia; nitrite;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ss; probe
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accumulation
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ID AAS11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT
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Best Local
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                     The invention relates to an oligonucleotide which includes at least 1 nucleic acid sequence which binds to at least 1 nuclear protein found in lung cells (e.g. the thyroid transcription factor 1, TTF-1, protein). The oligonucleotide can be expressed in lung cells via a vector and can be used to target therapeutic agents to kill lung or thyroid cancer cells. The oligonucleotide can be used to detect or diagnose lung or thyroid cancer. The oligonucleotides may be designed from the sequences of, for example, the promoters of lung-specific genes such as those encoding surfactant proteins. The present sequence is a mutated oligonucleotide from human surfactant protein B, SPB, promoter used in an EMSA (electrophoretic mobility shift assay) assay to confirm the presence of TTF-1 binding sites in the SPB promoter. The TTF-1 site has been mutated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 18
                                                                                                                                                                                                                                Oligonucleotide sequences which bind nuclear proteins and surfactants found in lung cells, useful for detecting cancers that originate in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human surfactant protein B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAS11828;
                                                                                                                                                                                                                                                                      WPI; 2001-513959/56.
                                                                                                                                                                                                                                                                                                  Bohinski RJ,
                                                                                                                                                                                                                                                                                                                                                               18-MAY-1994;
17-MAY-1995;
                                                                                                                                                                                                                                                                                                                                                                                                     26-MAY-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                              23-AUG-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                      US2001016352-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         mutation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Bcm; mutant.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Thyroid transcription
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; surfactant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             24-OCT-2001
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                                                                                                                                                                                           Example 5; Fig 34a; 76pp; English.
                                                                                                                                                                                                                     Lung.
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                                                                                                                                                                                                                                                                                                                          BOHINSKI R J.
WHITSETT J A.
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                                                                                                                                                                                                                                                                                                  Whitsett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Location/Qualifiers replace(10..13,CAAG) /*tag= a /note= "Mutates the TT
                                                                                                                                                                                                                                                                                                                                                               94US-00245356.
95US-00442809.
                                                                                                                                                                                                                                                                                                                                                                                                     9908-00320337.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              protein B; SPB; promoter; nuclear protein binding site;
tion factor; TTF-1; lung cancer; thyroid cancer; ds;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               entry
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  9 C; 2 G; 5 T; 0 U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               "Mutates the TTF-1
                                                                                                                                                                                                                                                                                                  JA;
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TTF-1 containing oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              42;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 binding
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                                                                                                                                                                                                                                     the
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Query Match Best Local S Matches 15

Similarity

80.0%; 93.8%; 6 C; 12

Score 14.4; Pred. No. 2.

6e+03

DB 4;

Length

30

Indels

0

Gape

0

15;

Conservative

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Sequence 30

BP; 6

P

G; 6 T; 0 U; 0 Other;

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                                                                                                                                                              Query Match
Best Local S
Matches 15
                                                                                                                                                                                                                                      This sequence represents a probe which was used in the detection of 16S rDNA sequence derived from an ammonia oxidising bacteria (AOB). The amplified sequence may be used in a composition which comprises an isolated bacterial strain that oxidizes ammonia to nitrite. The composition may be used for alleviating or preventing the accumulation of ammonia in a medium. The ammonia is reduced by at least 30% when compared with a level of ammonia that would exist in the absence of the bacterial strain. The composition is useful for alleviating or preventing the accumulation of ammonia in aqueous environment, e.g. a freshwater or
             17-SEP-2001
                                                                                                                                                                                                             Sequence
                                                                                                                                                                                                                                                                                                                                                                                           New composition comprising an isolated bacterial strain that oxidizes ammonia to nitrite, useful for alleviating or preventing the accumulation
                                    AAH42343;
                                                          AAH42343
                                                                                                                                                                                                                                                                                                                                                             Claim 10; Page
                                                                                                                                                                                                                                                                                                                                                                                    of ammonia in aqueous environment.
                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2004-304936/28.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    19-SEP-2002; 2002US-0386217P
19-SEP-2002; 2002US-0386218P
19-SEP-2002; 2002US-0386219P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Nitrosomonas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADM32724 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (AQUA-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    10-SEP-2003;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           01-APR-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO2004026772-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             aqueous
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Nitrosomonas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       17-JUN-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          16S rDNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             24
                                                                                                                   N
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                                                                                                                                                                         Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AQUARIA INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A; ammonia oxidising bacteria; AOB; ammonia; nitrite;
environment; freshwater; seawater; aquarium; ss; probe.
                                                          standard;
                                                                                                                                      CCCCCTCTTCTGGATAC 18
                                                                                                                                                                                                             18
                                                                                                                                                                                                                                  aquarium.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CCCCCTCTTCTGGATA
                                                                                                                                                              Conservative
                                                                                                                                                                                                           BP; 3 A; 9 C; 2 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2003WO-US028210
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      aestuarii.
            (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                aestuarii-like
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                             34; 98pp; English.
                                                          DNA;
                                                                                                                                                                        76.7%;
88.2%;
                                                          28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             9
                                                                                                                  18
                                                          BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ВP
                                                                                                                                                             0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                168
                                                                                                                                                                       Score 13.8; DB Pred. No. 5e+03;
                                                                                                                                                             Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            rDNA probe
                                                                                                                                                                                   DB 12;
                                                                                                                                                             2
                                                                                                                                                             Indels
                                                                                                                                                                                   Length
                                                                                                                                                                                    18;
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                                                                                                                                                            Gaps
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Homo sapiens Synthetic.

밁 8

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Matches
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Best Local
detection; cancer; 8q22.3; chromosome 8; human; EDD; tumour suppressor; cell cycle modulator; DNA repair; DNA damage; nuclear targeting protein; progesterone receptor; cytostatic; gene therapy; sequamous cell carcinoma; hepatocellular carcinoma; ovarian cancer; breast cancer; melanoma; head and neck cancer; adenocarcinoma; squamous lung cancer; gastrointestinal cancer; renal cell cancer; bladder cancer; prostate cancer; non-squamous carcinoma; glioblastoma; medullablastoma; carbonic anhydrase II; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present sequence represents a PCR primer for a cDNA fragment encoding an isoform of UDP-xylose;proteoglycan core protein beta-D-xylosylytransferase (XT). The XT enzyme occurs in at least two isoforms (XT-I) and (XT-II). XT is involved in the biosynthesis of glycosaminoglycans. XT polypeptides and polynucleotides may be used in the production of an agent (inhibitors and antagonists of XT) for the treatment of sclerotic diseases and chronic inflammatory joint diseases, or as a diagnostic marker. The XT DNA may be used a gene marker. Anti-XT antibodies are used as a diagnostic tool in a immunological assay for detection of a protein having XT activity
                                                                                                                                                                                                                                   Human carbonic anhydrase II forward PCR primer SEQ ID NO:31
                                                                                                                                                                                                                                                                                           03-JUN-2004
                                                                                                                                                                                                                                                                                                                                             ADL32573;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    UDP-xylose:proteoglycan core protein beta-D-xylosylytransferase and nucleic acids that encode it, useful for preventing, diagnosing and treating sclerotic diseases and chronic inflammatory joint diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI;
                                                                                                                                                                                                                                                                                                                                                                                              ADL32573
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     UDP-xylose:proteoglycan core protein beta-D-xylosylytransferase;
XT-I; XT-II; glycosaminoglycan; sclerotic disease; PCR primer;
chronic inflammatory joint disease; diagnostic marker; gene marker
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example
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01-DEC-2000; 2000EP-00126233
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (KLEE/) KLEESIEK K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                28-DEC-2000; 2000WO-EP013311.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2001-441872/47.
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                                                                                                                                                                                                                                                                                                                                                                                              standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CCCCCTCTTCTGGATAC 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CCCCCACTTCTGCATAC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              BP; 7 A; 2 C; 15 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
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                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  29; 80pp;
                                                                                                                                                                                                                                                                                                                                                                                           DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              76.7%;
88.2%;
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Pred.
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ase; diagnostic marker; gene marker; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           13.8; DB 4;
. No. 5.2e+03;
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05-SEP-2003; 2003WO-AU001164

18-MAR-2004 WO2004022750-A1

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cc activity, a protein having cell cycle modulatory activity, a protein associated with DNA repair or damage, a nuclear targeting protein, and a cc progesterone receptor protein or its portion, where the portion of the cc second polypeptide is sufficient to bind to the EDD polypeptide or the cc complex; (7) a method for determining a predisposition for disease, or cc disease state; (8) a method for determining a modulator of the activity, formation; (10) a method for determining a modulator of the activity, cc arid, ribozyme, peptide nucleic acid, (12) a pharmaceutical compatition acidl; (11) an antisense nucleic caid, ribozyme, pNA, interfering RNA or sirNA. EDD has cytostatic acid; and be used in gene therapy. The method and modulator are useful for creating a condition associated with sreast cancer, melanous cell carcinoma, hepatocellular carcinoma, ovarian cancer, cc lung cancer, gastrointestinal cancer (e.g. sguamous cell cancer, bladder cancer, prostate cancer, non-squamous carcinoma, plioblastoma and medullablatorom. The composition are useful to reducing the expression of EDD in a cell to inhibit cellular proliferation. The present sequence represents a PCR primer for human carbonic anhydrase II, which is used in the cc exemplification of the present invention.
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                               ð
                                                                       Query Match
Best Local S
Matches 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            linked to map position 8q22.3 of the human genome or its expression product in a sample of the subject, where an elevated level of the nucleic acid or polypeptide is indicative of cancer in the subject. Also described: (1) a method for diagnosing a cancer or predicting recurrence of a cancer in a subject comprising determining the level of mRNA or protein encoded by a nucleic acid as described above; (2) the isolated nucleic acid molecule for detecting cancer cell; (3) an isolated or recombinant protein complex; (4) an antibody that binds to the protein complex; (5) a kit for detecting or producing a protein complex, comprising an EDD polypeptide or a portion of an EDD polypeptide and a second polypeptides selected from a protein having tumour suppressor artivity a mortain having tumour suppressor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Detecting a cancer cell in a subject sample, also related to treatments, comprises determining the level of nucleic acid t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         05-SEP-2002;
07-NOV-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2004-248472/23
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                                                                                                                                                  Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              linked to map position 8q22.3 of the human genome or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              'n
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                                                                                            Similarity
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d R, O'brien
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQ ID NO 31; 331pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   invention describes a method
                                                                                                                                                    BP; 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2002AU-00951346.
2002US-0425218P.
                                                                                                                                                  ŗ
                                                                                            74.4%;
93.3%;
                                                                                                                                                    10 C; 3 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Henderson M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               scribes a method for detecting a cancer cell determining the level of nucleic acid that i
                                                                           0
                                                                           Pred. No. 8.1
); Mismatches
                                                                                          Score 13.4; DB 12
Pred. No. 8.1e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Clancy
                                                                                                              DB 12;
                                                                               Indels
                                                                                                            Length
                                                                                                                  22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   that
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04-DEC-2003

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RESULT 8
ADG31445/c
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Query Match
Best Local Similarity
Matches 15; Conserv
                                                                                                                                                                                                                                                                                                                                                                              The invention relates to a novel DNA encoding a weed controller metabolism protein. A protein of the invention has herbicide activity. The proteins and their encoded genes are useful e.g. in constructing new breeds of herbicide-resistant plants and also in developing various breeds of herbicide-resistant plants and also in developing various agrechemicals. The present sequence is used in the exemplification of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Weed controller metabolism proteins deactivating porphyrinogen oxidase (ppO)-inhibiting herbicides by N-demethylation and their genes, useful e.g. in constructing new breeds of herbicide-resistant plants.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  agrochemical;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Weed controller metabolism associated PCR primer SEQ ID NO:288
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                                                                                                                                                                                                                                                                                                                                               Sequence 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 19-OCT-2001;
07-JUN-2002;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              15-MAY-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               weed controller metabolism; weed; herbicide; herbicide-resistant plant;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             18-DEC-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADC36420;
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Disclosure; SEQ ID NO
                                                                              human; PCR; ss; primer; MN/CA IX; MN; oncogene; carbonic anhydrase 9; CA9; MN/CA9; carbonic anhydrase IX; CA IX; MN/G250; neoplastic tumour;
                                                                                                                                                                                       ADG31445 standard;
                                            Homo sapiens
                                                                   cytostatic;
                                                                                                                  PCR primer used to amplify human
                                                                                                                                         26-FEB-2004
                                                                                                                                                                                                                                                                                                                                                                       invention.
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                                                                                                                                                                                                                                                                                                                                               BP; 5 A; 4 C; 8 G; 3 T; 0 U; 0 Other;
                                                                    cancer;
                                                                                                                                                                                                                                                                                                 Conservative
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2002JP-00167239.
                                                                                                                                           (first
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                                                                                                                                                                                          DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             entry)
                                                                                                                                         entry)
                                                                                                                                                                                                                                                                                                           73.3%;
83.3%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                       812pp; Japanese.
                                                                    therapy; anti-tumour
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                                                                                                                                                                                                                                                                                                               Score 13.2; DB 10;
Pred. No. 1e+04;
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                                                                                                                    c-erbB2 DNA
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                                                                                                                                                                                                                                                                                                             1e+04;
                                                                                                                    SeqID
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RESULT 9
ADK41856/c
ID ADK418
XX ADK418
AC ADK418
XX Human
DT O6-MAY
XX Carbon
KW Carbon
KW renal
XX Homo 8
XX Homo 9
XX Homo 
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Best Local S
Matches 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         mice). Specifically, this antibody is directed towards the MN gene, a cellular oncogene known alternatively as carbonic anhydrase 9, CA9 or MN/CA9, which encodes the MN protein that is also known as the MN/CA IX isoenzyme, carbonic anhydrase IX. CA IX or the MN/G250 protein. The present invention describes the generation of this monoclonal antibody, and immunoreactive fragments thereof, which are directed against non-immunodominant epitopes on the CA IX extracellular domain. As such, this antibody can be useful diagnostically as a marker for preneoplastic/ neoplastic tumours, immunodetection methods and immunotargeting approaches. Accordingly, compositions exhibit cytostatic activity and are useful in the diagnosis, prognosis and treatment of various cancers including breast, bladder or lung cancer, in tumour therapy and in antitumour vaccination. This oligonucleotide sequence is an PCR primer used
                                                                                                                                                                                                                                                                                                  carbonic anhydrase IX; CA IX; precancerous cell; MN; cancerous cell; human; vertebrate; cytostatic; vaccine; gene therapy; renal cell carcinoma; breast cancer; colorectal cancer; PCR; primer;
     23-MAY-2002;
05-DEC-2002;
                                                                                    22-FEB-2003; 2003WO-US005137.
                                                                                                                                                                                                                                                                                                                                                                                                                      Human MN gene-related PCR primer SeqID109
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                06-MAY-2004
                                                                                                                                                                                                  WO2004005348-A1
                                                                                                                                                                                                                                                   Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADK41856
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 27 BP; 6 A; 6 C; 10 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    This invention relates to a novel monoclonal antibody identified as the MN/CA IX specific antibody prepared in knockout mice (CA IX deficient
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New monoclonal antibody generated from MN/CA IX-deficient mice, where antibody binds specifically to human tumor-associated cell adhesion protein MN/CA IX or polypeptide, useful for diagnosing, prognosing or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 11; SEQ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CCCTCCTCATCTGGAAAC 2
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  2002US-0383068P.
2002US-0431499P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
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2002US-0431499P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           human breast tumour marker,
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Pred.
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No. 1e+04;
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RESULT 10
ADC36414
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Best Local S
Matches 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           This invention relates to a novel soluble form of the carbonic anhydrase IX (CA IX) (or MN) protein or CA IX polypeptide which is released from precancerous and/or cancerous cells of a vertebrate into a body fluid. The invention may be useful for the development of compounds with a cytostatic activity or a vaccine whilst the disclosed sequences may be used for gene therapy. The protein and method are useful for screening, diagnosing or prognosing diseases associated with abnormal expression of carbonic anhydrase IX protein, such as precancerous and cancerous diseases like renal cell carcinoma, breast cancer or colorectal cancer. The monoclonal antibody may also be used for treating or preventing precancerous and cancerous diseases. The present sequence is that of a PCR primer which was used in the exemplification of the invention.
                                                                                    Nakajima
                                                                                                                                   19-OCT-2001;
07-JUN-2002;
                                                                                                                                                                     17-OCT-2002; 2002WO-JP010789
                                                                                                                                                                                                                                                                         weed controller metabolism; weed; herbicide; herbicide-resistant plant; agrochemical; ss; PCR; primer.
                                                                                                                                                                                                                                                                                                                Weed
                                                                                                                                                                                                                          WO2003040370-A1.
                                                                                                                                                                                                                                                 Synthetic
                                                                                                                                                                                                                                                                                                                                                                 ADC36414;
                                                                                                                                                                                                                                                                                                                                                                                          ADC36414 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New soluble form of the carbonic anhydrase IX (CA IX) protein for screening, diagnosing or prognosing diseases associated with abnormal expression of CA IX protein, e.g. renal cell carcinoma, breast cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (FARB )
(VIRO-)
                                                                                                                                                                                                                                                                                                                                        18-DEC-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           expression colorectal
                                                                                                             (SUMO)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Zavada J,
                                                                                                                                                                                                                                                                                                            controller metabolism associated PCR primer SEQ ID NO:282.
                                                                                                                                                                                                                                                                                                                                                                                                                                                         19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               μ
                                                                                                             SUMITOMO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Similarity
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INST VIROLOGY
                                                                                  Ħ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CCCCCCTCTTCTGGATAC 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  diagnosing or prognosing disc
of CA IX protein, e.g. renal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BP; 6 A; 6 C; 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                    Mukumoto
                                                                                                                                   2001JP-00321307
2002JP-00167239
                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ID NO 113;
                                                                                                           CHEM CO
                                                                                                                                                                                                                                                                                                                                                                                          DNA; 28
                                                                              T
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   83.3%;
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                                                                                    Takaishi
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Pastorek J,
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Pred.
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5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              3 13.2; UB 1.
No. 1e+04;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 12;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    carcinoma, breast cancer
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Weed controller metabolism proteins deactivating porphyrinogen oxidase (PPO)-inhibiting herbicides by N-demethylation and their genes, useful e.g. in constructing new breeds of herbicide-resistant plants.

2003-523102/49

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RESULT 11
ACA54549/c
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The invention relates to an isolated nucleic acid comprising any one of the 6213 antisense sequences given in the specification where expression of the nucleic acid inhibits proliferation of a cell. Also included are: (1) a vector comprising a promoter operably linked to the nucleic acid encoding a polypeptide whose expression is inhibited by the antisense nucleic acid; (2) a host cell containing the vector; (3) an isolated polypeptide or its fragment whose expression is inhibited by the antisense nucleic acid; (4) an antibody capable of specifically binding the polypeptide; (5) producing the polypeptide; (6) inhibiting cellular proliferation or the activity of a gene in an operon required for proliferation; (7) identifying a compound that influences the activity of the gene product or that has an activity against a biological pathway
                                                                                                                                                                                                                                                                                                                                                            21-MAR-2001;
06-SEP-2001;
25-OCT-2001;
08-PEB-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 grug
                                                                                                                                                                                                    New antisense nucleic acids, useful for identifying proteins or i for homologous nucleic acids required for cellular proliferation isolate candidate molecules for rational drug discovery programs
                                                                                                                                                                                                                                                                                                                                                                                                                              21-MAR-2002; 2002WO-US009107
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   27-OCT-2003
19-JUN-2003
                                                                                                                                                                          Example
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WO200277183-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Antisense;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Prokaryotic
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                                                                                                                                                                                                                                                        2003-029926/02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   design;
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                                                                                                                                                                                                                                                                                                                       ELITRA PHARM
                                                                                                                                                                          21; Page 1574; 1766pp; English.
                                                                                                                                                                                                                                                                                  Zamudio
Trawick
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CCACCGTCTCCTGGATAC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ss; prokaryotic essential gene; cell proliferation;
n; primer; PCR.
                                                                                                                                                                                                                                                                                                                                              ; 2001US-00815242.
; 2001US-00948993.
; 2001US-0342923P.
; 2002US-00072851.
; 2002US-0362699P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          folA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (revised)
(first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          gene PCR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DNA;
                                                                                                                                                                                                                                                                                                                        INC.
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Carr (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   30
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   28
                                                                                                                                                                                                                                                                                  Haselbeck R, Yamamoto R,
                                                                                                                                                                                                                                                                                  Ohlsen
Forsyth
                                                                                                                                                                                                                                                                                    ₽°
                                                                                                                                                                                                                                                                                    Zyskind
Xu HH;
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(ELIT-)

ELITRA PHARM INC

09-FEB-2001; 2001US-0267636P

08-FEB-2002;

2002WO-US003987

Carr Go, Zyskind JW

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ž Виввеу Н;

Wall

Foulkes GJ, D, Trawick

Zamudio C, F JD, Yamamoto

Haselbeck R, ያ ነ

'n Ohlsen

Ď,

WPI;

2003-093128/08

Identifying the target of a compound which interproliferation, comprises contacting a culture or underexpress a gene product with the above

inhibits cellular ure of strains that overexpress ove compound, and identifying

Example

16;

Page

315;

640pp;

English

gene

product.

This invention describes a novel method for identifying gene products compounds inhibiting proliferation of an organism act. The method comprises obtaining a culture of strains overexpressing a different product for proliferation of the organism, contacting the culture with compound to inhibit proliferation of strains that do not overexpress the

ducts on method

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                                                                                                                                                                                                                                                                                                                                                                     cc required for proliferation, or that inhibits cellular proliferation; (8) cidentifying a gene required for cellular proliferation or the biological companies on which the test compound that inhibits proliferation of an cc compound's activity; (11) a culture comprising strains in which the gene product is overexpressed or underexpressed; (12) determining the extent cc which each of the strains is present in a culture or collection of strains; or (13) identifying the target of a compound that inhibits the compound that inhibits 
                                                                                                                                                       Query Match
Best Local
                                                                                                                         Matches
                                                                                                                                                                                                                                                         Sequence
                                                                                                                                                                                                                                                                                                                  allows growth in the presence elevated (Updated on 27-OCT-2003 to standardise
21
                                                         \vdash
                                                                                                                         l Similarity
                                                                                                                                                                                                                                                         30
                                 CCCCCCTCTTCTGGATAC
                                                                                                                                                                                                                                                         BP; 11 A; 3 C; 10 G; 6 T;
                                                                                                                            Conservative
                                                                                                                                                       73.3%;
                                                            18
4
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Pred.
                                                                                                                                Mismatches
                                                                                                                                                              13.2;
No. 1e
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OS field)
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CCCCCCTCTTCTGGATAC 18

Query Best L Matches

Local Match

Similarity

73.3**%**; 83.3**%**;

Conservative

0;

Score 13.2; DB Pred. No. 1e+04; 0; Mismatches

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0

Gaps

0

Length Indels

28;

Sequence 28 BP; 5

A; 11 C;

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T; 0 ů,

0 Other DB 10;

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Disclosure;

SEQ

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282;

812pp; Japanese

The invention relates to a novel DNA encoding a weed controller metabolism protein. A protein of the invention has herbicide activity. The proteins and their encoded genes are useful e.g. in constructing new breeds of herbicide-resistant plants and also in developing various agrochemicals. The present sequence is used in the exemplification of the

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ABX13526/c
ID ABX13526
                                                       Cellular proliferation; inhibitor; target; primer; PCR; trimethroprim; 88.
                                                                                                                                    04-JUN-2003
                                                                                                                                                                 ABX13526;
WO200286097-A2
                           Staphylococcus
                                                                                                       aureus
                                                                                                                                                                                                 standard; DNA; 30
                                                                                                      fola PCR primer SEQ ID 15798
                                                                                                                                      (first entry
                              aureus
                                                                                                                                                                                                  ₽₽
                                                                            folA;
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product and identifying the product overexpressed in a strain that proliferated more rapidly. The method is useful in identifying the ta of a compound which reduces the activity or level of gene products required for cellular proliferation. The method may also be used for identifying the therapeutic compounds that act on the novel targets. sequence represents a PCR primer used to amplify the Staphylococcus aureus folk gene which is inhibited by trimethroprim the target Thie

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Sequence 30 BP; 11 A; 3 C; 10 G; 6 T; 0 U; 0 Other;

δ Query Match Best Local S Matches 15 15; ,, Similarity CCCCCCTCTTCTGGATAC Conservative 73.3%; 83.3%; 18 0 Score 13.2; Pred. No. 1 Mismatches le+04; 멂 80 ų. Length Indels 0 Gaps

RESULT 13 ADH30251 밁 21 CCTCCTTCTTCTGGATCC

4

ADH30251 standard; DNA; 17 ВP

ADH30251; (first entry)

Human monocyte-macrophage cell surface apoB48 receptor primer #5

antilipemic; gene therapy; monocyte-macrophage cell surface apoB48 receptor; apoB48R; pattern B phenotype; familial combined hyperlipidaemia; hypercholesterolaemia; hypertriglyceridaemia; low plasma high-density lipoprotein level; human; primer;

Homo sapiens.

US2003208060-A1

06-NOV-2003

12-JUN-2003; 2003US-00459876

31-MAY-2000; 06-AUG-1998; 98US-00130242. 2000US-00583610.

(GIAN/) (BRAD/) GIANTURCO BRADLEY W ≯ S Ξ

Gianturco

SH,

Bradley

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2003-864799/80

New isolated DNA molecule encoding a monocyte-macrophage cell surface apoB48 receptor protein (apoB48R), useful in preparing a composition treating e.g., familial or non-familial hypercholesterolemia. for

Example 23; SEQ ID NO 17; 77pp; English.

The invention describes a new isolated DNA molecule, encoding a monocyte-macrophage cell surface apo848 receptor protein (apo848R) comprising a fully defined 1088 amino acid sequence. The DNA molecule is useful in preparing a composition for treating Pattern B phenotype, familial combined hyperlipidaemia, familial or non-familial hypercholesterolaemia, hypertriglyceridaemia or low plasma high-density lipoprotein levels. This sequence represents a primer used in the isolation of human monocytemacrophage cell surface apo848 receptor DNA.

Sequence 17 B₽; 2 A; 6 C; 4 G; 5 T; 0 U; 0 Other;

Query Match Best Local S Matches 14 Similarity Conservative 71.1%; 0, Score 12.8; Pred. No. 1 Mismatches .6e+04 DB 10; Indels Length 17; <u>;</u> Gaps

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      S
         ب
         CCCCCCTCTTCTGGAT
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RESULT 14

AD000840 standard; DNA; 17 ₽P

ADC00840

12-AUG-2004 (first entry)

0

Human monocyte-macrophage cell surface apoB48R cDNA PCR primer #6.

apolipoprotein B 48 receptor; apoB48R; PCR; B8; monocyte; mareticulcendothelial cell; monocyte adhesion; cardiovascular apoB-mediated cellular uptake; plasma chylonicron; Human; monocyte-macrophage cell ll surface apolipoprotein apoB48R; PCR; ss; monocyt W 48 receptor; macrophage; disease;

hypertriglyceridaemic triglyceride-rich lipoprotein;

hypertriglyceridaemia; primer. ApoE independent pathway; lipoprotein lipase independent pathway; heparin sulfate proteoglycan independent pathway; foam cell formation; ApoB independent pathway;

Homo sapiens

US6740735-B1

25-MAY-2004

31-MAY-2000; 2000US-00583610

06-AUG-1998; 98US-00130242

(GIAN/) GIANTURCO S (BRAD/) BRADLEY W A. Ξ

Gianturco SH, Bradley Æ

WPI; 2004-387250/36.

New isolated monocyte-macrophage cell surface apoB48 receptor useful for cell-specific delivery of therapeutic compounds to monocytes or macrophages, or for evaluating an individual at cardiovascular disease. risk for human

Example 23; SEQ ID NO 17; 72pp; English

ADOOO840

ADOOO840

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AC

ADOO

XX

ADOO

AC

ADOO

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ADOO

DT 12-A

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ADOO

DT 12-A

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Huma

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Huma

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A cc delivery of therapeutic compounds to human moncytes, macrophages or cother reticuloendothelial cells that express the receptor, a method of cinhibiting foam cell formation and increased monocyte adhesion to cendothelial cells and a method of evaluating an individual at risk of a cardiovascular disease. The monocyte-macrophage cell surface apoB48 creceptor protein is useful in the apoB-mediated cellular uptake of plasma cc hylomicrons and remnants and hypertriglyceridaemic triglyceride-rich cc lipoproteins in an ApoB independent pathway, a lipoprotein lipase cc independent pathway or a heparin sulfate proteoglycan independent cc pathway. It is also useful for the cell-specific delivery of therapeutic compounds to human monocytes, macrophages or other reticuloendothelial cells that express the receptor, for inhibiting foam cell formation and conceased monocyte adhesion to endothelial cells and for evaluating an individual at risk of a cardiovascular disease such as compounded to human monocytes achieved as a popular disease such as compounded to human monocytes achieved achieved as a popular disease such as compounded to human monocytes achieved achieved as a popular disease such as compounded to human monocytes achieved achieved achieved as cardiovascular disease such as compounded to human monocytes achieved achieved achieved achieved as cardiovascular disease such as compounded to human monocytes achieved a The invention relates to the human monocyte-macrophage cell surface apolipoprotein B 48 receptor (apoB48R) protein and the nucleic acid encoding it. The invention also relates to a method of cell-specific

Sequence 17 BP; N Α σ C; <u>ن</u> σı Η. 0 ς, 0 Other

Query Match Best Local Similarity 71.1%; 87.5%; Score Pred. No. 1. 8; DB 12; 1.6e+04; Length 17;

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RESULT 15
ADNO8161/c
ID ADNO81
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                                                                                                                               The invention relates to a composition that contains, as active agent, an ccid, polypeptide or antisense nucleic acid, polypeptide or antisense nucleic acid, antibody against EG-VEGF or cits receptor (EG-VEGFR), or EG-VEGFR antisense nucleic acid, is useful cor the treatment or prevention of endometrial diseases. Also included cor are a method for detecting uterine receptivity (by determining the amount cor feg-VEGF polypeptide and/or nucleic acid, using the new composition) or a test system for identifying antagonists of EG-VEGF-R. Also disclosed as new is a splice variant of the human GPR73a receptor. It contains an additional exon which includes a stop codon, so produces a truncated cor protein that can not mediate signal transduction, i.e. it functions as a composition that can not mediate signal transduction, i.e. it functions as a composition that can not mediate signal transduction, i.e. it functions as a composition that can not mediate signal transduction, i.e. it functions as a composition of edges to the composition of the composition of the best time for composition of eggs fertilised in vitro. Also the test system that composition is used to screen for composition and the express the EG-VEGF receptor is used to screen for composition of edges fertilised in vitro. Also the test system that composition is used to screen for composition and the isolation of the EG-VEGF cDNA.
                                 Query Match
Best Local S
Matches 14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human; 88; PCR; endocrine gland vascular endothelial growth factor; EG-VEGF; EG-VEGF receptor; endometrial disease; uterine receptivity; endometriosis; endometrial carcinoma; dysfunctional bleeding; gene therapy; primer; RT-PCR; reverse transcriptase PCR; S9-RNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADNO8161 standard; DNA; 18 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 1; Page 5; 13pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Treatment and prevention carcinoma, by inhibiting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2004-134408/14.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Haendler B, Hess-Stumpp
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        26-JUN-2002; 2002DE-01029379
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      26-JUN-2002; 2002DE-01029379.
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87.5%;
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Listing first 45 summaries
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Maximum DB seq length: 2000000000
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1: gb_ba:*
2: gb_in:*
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Gapop 10.0 , Gapext 1.0
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Copyright (c) 1993 - 2006 Biocceleration Ltd.
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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AY283110	AY123812	AY123803	AY856079	AF386754	AF386755	NTV16SRRNA	NS16SAV	NTV16RRNAA	NSAF16S	NSP298724	AF408634	AY921481	AF134442	AF134441	AX316099	CS089161	CQ796915	ID
AY283110 Unculture	AY123812 Nitrososp	AY123803 Nitrosovi	AY856079 Nitrososp	AP386754 Nitrososp	AF386755 Nitrososp	M96397 Nitrosovibr	Y10127 Nitrosospir	M96404 Nitrosovibr	X84658 Nitrosospir	AJ298724 Nitrososp	AF408634 Nitrososp	AY921481 Unculture	AF134442 Unculture	AF134441 Unculture	AX316099 Sequence	CS089161 Sequence	CQ796915 Sequence	Description

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1493	1492	1492	1486	1474	1458	1458	1458	1458	1457	1448	1447	1445	1442	1434	1399	1321	1319	1294	1285	1217	1207	1207	1207	1206	692	569
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AY218755	AY218730	AY218727	AF386756	NSS16SR	AX316094	CS089156	CQ796910	AY684260	AF353162	NTR16SRRNA	AF080256	AF353157	AF353158	NTV16RRNAB	NSL11516S	NSAJ5545	NSAJ5544	NSAJ5543	USO390469	AY726896	AY727033	NTE298746	NSP298747	AY727034	AY534019	AJ704699
AY218755 Unculture	AY218730 Unculture		AF386756 Nitrososp	L35505 Nitrosospir		CS089156 Sequence			AF353162 Nitrososp				AF353158 Nitrososp	M96405 Nitrosovibr	₽-	AJ005545 Nitrososp	AJ005544 Nitrososp	AJ005543 Nitrososp	AJ390469 Unculture		AY727033 Nitrososp	AJ298746 Nitrososp	AJ298747 Nitrososp	AY727034 Nitrosos	AY534019 Unculture	AJ704699 Unculture

ALIGNMENTS

RESULT 2 CS089161 LOCUS DEPINITION ACCESSION VERSION VERSION KEYMORDS SOURCE	Query Match Best Local Matches 1 Qy 1 Db 1	SOURCES SOURCES SOURCES ORGANISM REFERENCE AUTHORS TITLE JOURNAL FEATURES BOUICE ORIGIN	RESULT 1 CQ796915 LOCUS DEFINITION ACCESSION VERSION
CS089161 19 bp DNA Sequence 8 from Patent EP1502948. CS089161.1 GI:66714445 synthetic construct	100.0%; Score 19; DB 6; Similarity 100.0%; Pred. No. 20; 19; Conservative 0; Mismatches 0; 1 TCCCCCACTCGAAGATACG 19	synthetic construct synthetic construct other sequences; artificial sequences. Hovanec,T.A. Ammonia-oxidizing bacteria and methods of thesame Patent: WO 2004026772-A 8 01-APR-2004; Aquaria Inc. (US) Location/Qualifiers 119 /organism="synthetic construct" /mol_type="nuassigned DNA" /db_xref="taxon:32630" /note="Oligonucleotide probe"	CQ796915 I9 bp DNA Sequence 8 from Patent WO2004026772. CQ796915 CQ796915.1 GI:46408541
linear	Length 19; Indels	using and	linear
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ACCESSION
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uncultured beta proteobacterium b4
Bacteria, Proteobacteria, Betaproteobacteria, Nitrosomonadales,
Nitrosomonadaceae, environmental samples.
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McCarthy,A.
Phylogehetic differentiation of two closely related Nitrosomonas
spp. That inhabit different sediment environments in an
                                                                                                                                                                                                 Uncultured beta
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Sequence 8 from
AX316099
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Patent: WO 0190312-A 8 29-NOV-2001;
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Patent: EP 1502948-A 8 02-FEB-2005;
Aquaria Inc. (US)
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                                                    Whitby, C.B., Saunders, J.R.,
                                                                                                                                                                                    sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            other sequences; artificial sequences
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/db_xref="taxon:32630"
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|mol_type="unassigned DNA"
|db_xref="taxon:32630"
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Patent W00190312.
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                                                    Rodriguez, J., Pickup, R.W. and
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ch 100.0%;
l Similarity 100.0%;
l9; Conservative 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               sequence.
AF134442
                                                                                                                                                                                                                                                                                    2 (bases 1 to 455)
Whitby, C.W., Saunders, J.R.,
McCarthy, A.J.
                                                                                                                                                                                                                                                                                                                                   Appl. Env
10543796
                                                                                                                                                                                                                                                                                                                                                                                                                                                 uncultured beta proteobacterium b5
uncultured beta proteobacterium b5
Bacteria; Proteobacteria; Betaproteobacteria; Nitrosomonadales;
Nitrosomonadaceae; environmental samples.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 \vdash
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                19;
                                                                                                                                                                                                                                        Submitted (12-MAR-1999) School of Biological Sciences, of Liverpool, Crown Street, Liverpool, Merseyside L69
                                                                                                                                                                                                                                                                                                                                                 Phylogenetic differentiation of two closely related Nitrosomonas spp. That inhabit different sediment environments in an oligotrophic freshwater lake Appl. Environ. Microbiol. 65 (11), 4855-4862 (1999)
                                                                                                                                                                                                                                                                                                                                                                                                             Whitby, C.B., Saunders, J.R., Rodriguez, J., Pickup, R.W. McCarthy, A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ENV
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Submitted (12-MAR-1999) School of Biological Sciences, University of Liverpool, Crown Street, Liverpool, Merseyside L69 72B, UK
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                oligotrophic freshwater lake Appl. Environ. Microbiol. 65 10543796
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Similarity 100.0%; Pred. No. 18;
19; Conservative 0; Mismatches
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/mol_type="rRNA"
/db_xref="taxon:93622"
/chromosome="1"
                                                                                                      /environmental_sample
/note="similar to Escherichia
                                                                                                                                                 /db_xref="taxon:93626"
/chromosome="1"
                                                                                                                                                                               /organism="uncultured beta proteobacterium
mol_type="rRNA"
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/note="similar to Escherichia
                                                                           product="16S ribosomal RNA"
                                                                                                                                      clone="b5"
                                                                                                                                                                                                                        ocation/Qualifiers
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              Score 19;
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red. No. 18;
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                            Length 455;
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AF408634/c
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                                                                                                                                                                                                                                                                                                                                                                            365
                                     2 (bases 1 to 983)
Norton, J.M. and Klotz, M.G.
Direct Submission
Submitted (10-AUG-2001) plant, (
University, 4820 Old Main Hill,
                                                                                                                                                                        Nitrosospira multiformis
Bacteria; Proteobacteria; Betaproteobacteria;
Nitrosomonadaceae; Nitrosospira.
                                                                                                                                                                                                                                                               Bequence.
AF408634
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              28
                                                                                                                                   1 (bases 1 to 983)
Norton, J.M., Alzerreca, J.J.,
Diversity of the amo operon
                                                                                                                                                                                                                                                                                      AF408634
Nitrosospira
                                                                                                                                                                                                                                                                                                                                                                                                                                 19;
                                                                                                                                                                                                                     Nitrosospira multiformis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Submitted (06-FEB-2005) School Seoul National University, San
                                                                                                                          bacteria
                                                                                                                                                                                                                                                   AF408634.1
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Ka, J.-O., Jung, M.-K., Kim, M.-S. and Ahn, J.-H.
Direct Submission
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Unpublished
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 (bases 1 to 493)
Ka,J.-O., Jung,M.-K., Kim,M.-S. and Ahn,J.-H.
Changes in bacterial community in rice field soils
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Bacteria; environmental samples.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         uncultured bacterium
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organism="Nitrosospira multiformis"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /environmental_sample
complement(<1._.>493)
/product="168 ribosomal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Korea
                        ocation/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /organism="uncultured bacterium"
/mal_type="genomic DNA"
/isolation_source="rice field soil"
/db_xref="Eaxon:77133"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              clone="AP5"
                                                                                                                                                                                                                                                   GI:18139825
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multiformis strain 24C
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tive 0; Mismatches
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clone
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AP5
                                        Soils and Biometeorology,
Logan, UT 84322, USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  of Applied Biology and Chemistry, 56-1 Silimdong Kwanakgu, Seoul
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RNA"
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16S ribosomal
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RNA gene, partial
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            Nitrosospira sp.
                                   X84658.1 GI:1236774
16S ribosomal RNA.
                                                                        NSAF16S
Nitrosospira
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Nitrosospira sp. 24C
Bacteria; Proteobacteria; Betaproteobacteria; Nitrosomonadales;
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                                                               X84658
                                                                                                                                                                                                                                                                                                                                                                                                                Submitted (30-AUG-2000) Aakra A., Le
Technology, Agricultural University
N-1432 Aas, NORWAY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Detailed phylogeny of ammonia-oxidizing bacteria determined by sequences and DNA homology values
Int. J. Syst. Evol. Microbiol. 51 (Pt 6), 2021-2030 (2001)
Bacteria; Proteobacteria; Betaproteobacteria; Nitrosomonadales;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Aakra,A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AJ298724.1 GI:17932725
16S ribosomal RNA; 16S
                                                                                                                                                                                                              Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Nitrosomonadaceae; Nitrosospira.
                                                                                                                                                                       TCCCCCACTCGAAGATACG 19
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                                                                                                                                                                                                  Conservative
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llarity 100.0%;
Conservative 0
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/product="16S ribosomal RNA"
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                                                                                                                                                                                                                                                                                                  gene="16S rRNA"
                                                                                                                                                                                                                                                                                                                                        /isolate="24C"
/db_xref="taxon:136540"
                                                                                                                                                                                                                                                                                                                                                              /organism="Nitrosospira
/mol type="genomic DNA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /db_xref="taxon:1231"
<1. .>983
                                                                                                                                                                                                                                                                                                                           country="USA:Minnesota"
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/strain="24C"
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ty of Norway, P.O. Box 5051,
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Nitrosospira sp. NpAV pa
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Y10127.2 GI:27262953
16S rippsomal RNA; 16S r
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1 (bases 1 to 1378)
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16S ribosomal RNA.
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M96404
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                                                                                                                                                                                                                                                                                                                                              The phylogeny of autotrophic ammonia-oxidizing bacteria as determined by analysis of 16S ribosomal RNA gene sequences J. Gen. Microbiol. 139 (Pt 6), 1147-1153 (1993)
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/mol_type="unassigned DNA"
/db_xref="taxon:1233"
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McCaig, A.E.

Direct Submission

Submitted (08-NOV-2002) A.E. McCaig, Institute of Medical Sciences, Submitted (08-NOV-2002) M.E. McCaig, Institute of Medical Sciences, Molecular and Cell Biology, University of Aberdeen, Foresterhill, Molecular and Cell Biology, University of Aberdeen, Foresterhill, Molecular and Cell Biology, University of Aberdeen, Foresterhill, Aberdeen, AB25 2ZD, UK

On Dec 19, 2002 this sequence version replaced g1:1771383.

Location/Qualifiers
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Submitted (18-DEC-1996) A.B. McCaig, Institute of Medical Sciences,
Molecular and Cell Biology, University of Aberdeen, Foresterhill,
Aberdeen, AB25 ZZD, UK
revised by [4]
4 (bases 1 to 1388)
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Nitrosospira sp. NpAV
Bacteria; Proteobacteria; Betaproteobacteria; Nitrosomonadales;
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                                                                                                                                                                                                                                                           M96397.1 GI:150441
16S ribosomal RNA.
Nitrosovibrio tenuis
                                                                                                  The phylogeny of autotrophic ammonia-oxidizing bacteria as determined by analysis of 16S ribosomal RNA gene sequences J. Gen. Microbiol. 139 (Pt 6), 1147-1153 (1993)
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Nitrosovibrio tenuis 16S
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Nitrosomonadaceae; Nitrosovibrio.
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                                                                    Original
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                                                    source text: Nitrosovibrio tenuis Location/Qualifiers
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/mol_type="genomic DNA"
/strain="NpAV"
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Nitrosospira sp. BF16c46
Bacteria; Proteobacteria; Betaproteobacteria; Nitrosomonadales;
Nitrosomonadaceae; Nitrosospira; environmental samples.
                               1 (bases 1 to 1486)
Burrell, P.C., Phalen, C.M. a
Identification of bacteria
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Nitrosospira sp. BF16c46 16S
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Burrell,P.C., Phalen,C.M. and Hovanec,T.A.
Direct Submission
Submitted (30-MAY-2001) Aquatic Research Laboratory,
Group, 6100 Condor Dr, Moorpark, CA 93021, USA
Location/Qualifiers
                                                                                                                                        AF386754.1
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Nitrosospira sp. R5c20
Bacteria; Proteobacteria;
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Nitrosomonadaceae; Nitrosospira; environmental samples.
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/isolation source="mammonia-oxidizing
from freshwater aquaria enrichments"
/db_xref="taxon:180514"
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                                                                                                                                          GI:17864831
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sp. R5c20 16S ribosomal RNA
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Pred. No. 18;
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Pred. No.
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                            and Hovanec,T.A.
a responsible for ammonia oxidation
            67
         (12), 5791-5800 (2001)
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Nitrosospira sp. En13
Bacteria; Proteobacteria; Betaproteobacteria; Nitrosomonadales;
Nitrosomonadaceae; Nitrosospira.
1 (bases 1 to 1495)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Nitrosospira sp. Enl3 16S ribosomal
                                                                                                                                                                                                                                  Direct Submission
Submitted (14-DEC-2004) School of Biological Sciences, University of Aberdeen, Cruickshank Building, St Machar Drive, Aberdeen AB24
                                                                                                                                                                                                                                                                                  2 (bases 1 to 1495)
Shaw,L.J., Nicol,G.W.,
                                                                                                                                                                                                                                                                                                               Nitrous oxide production and nitrifier denitrification betaproteobacterial ammonia oxidising bacteria Unpublished
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                                                                                                                                                                                                                                                                             Baggs, E.M.
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Burrell, P.C., Phalen, C.M. and Hovanec, T.A.
Direct Submission
                                                              Similarity
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TCCCCCACTCGAAGATACG
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                                                   Conservative
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                                                                                                                 /product="16S ribosomal
                                                                                                                               /strain="En13"
/db_xref="taxon:308023"
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                                                                                                                                                                                               Location/Qualifiers
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mol type="genomic DNA"

/isolation_source="mamonia-oxidizing bacteria isolated from freshwater aquaria enrichments"

/db_xref="taxon:180513"
                                                                                                                                                                   organism="Nitrosospira/mol_type="genomic DNA"
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1: /SIDS5/ptodata/1/pubpna/US08_NEW_PUB.seq:*
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9: /SIDS5/ptodata/1/pubpna/US10_NEW_PUB.seq:*
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11; /SIDS5/ptodata/1/pubpna/US11_NEW_PUB.seq:*
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Copyright (c) 1993 - 2006 Biocceleration Ltd.
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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19 19 19	19 19 19	125 125 127 128	Length 25 25
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US-11-101-244-869498 US-11-101-244-869507 US-11-101-244-1029568	US-11-101-244-603439 US-11-101-244-815438 US-11-101-244-815537 US-11-101-244-817168	US-10-03-784-1230837 US-10-310-914A-224598 US-10-934-048A-63320 US-11-121-849-325988 US-10-310-914A-224589 US-10-310-914A-493308 US-10-310-914A-493308 US-10-310-914A-1226662	ID US-10-310-914A-1061199 US-11-121-849-112898
Sequence 869498, Sequence 869507, Sequence 1029568,	Sequence 603439, Sequence 815438, Sequence 815537, Sequence 817168,	Sequence 1230637, Sequence 1230837, Sequence 224598, Sequence 68320, A Sequence 32598, Sequence 224589, Sequence 23172, Sequence 1226662,	Description Sequence 1061199, Sequence 112898,

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Sequence 230052,	Sequence 596021,			Sequence 960621,	Sequence 541699,	Sequence 1518708,	Sequence 1425762,		æ								Sequence 815537,		Sequence 603439,			Sequence 1404224,	æ			Seguence 1059817,

ALIGNMENTS

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RESULT 1
US-10-310-914A-1061199/c
US-10-310-914A-1061199, Application US/10310914A

Publication No. US20060033322A1

[GENERAL INFORMATION: Bentwich, Isaac
| APPLICANT: John Palma

US-11-121-0914A-1061199

Query Match
| ORANISM: Human
| John Palma
| ORANISM: Human
| ORANISM: Human
| ORANISM: Human
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| ORANISM: John Palma
| ORANISM: John Palma
| ORANISM: John Palma
| Sequence 112898, Application US/11121849
| PHILEARTION: John Palma
| TITLE OF INVENTION: Methods of Genetic Analysis of Formalin Fixed Paraffin Embedded
| TITLE OF INVENTION: Methods of Genetic Analysis of Formalin Fixed Paraffin Embedded
| ORANISM: John Palma
| ORANISM: John Palma
| TITLE OF INVENTION: Methods of Genetic Analysis of Formalin Fixed Paraffin Embedded
| ORANISM: John Palma
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GENERAL INFORMATIÓN:

APPLICANT: Charmacon, Inc.

APPLICANT: Khvorova, Anastasia

APPLICANT: Khvorova, Anastasia

APPLICANT: Keynolds, Angela

APPLICANT: Leake, Devin

APPLICANT: Marshall, William

APPLICANT: Scaringe, Stephen

ITITE OF INVENTION: Functional and Hyperfunctional sirNA

FILIR REFERENCE: 13499US

CURRENT APPLICATION NUMBER: US/11/101,244

CURRENT FILING DATE: 2005-04-07

PRIOR APPLICATION NUMBER: 60/502,050

PRIOR FILING DATE: 2003-09-10

PRIOR PILING DATE: 2003-09-10

PRIOR APPLICATION NUMBER: 60/426,137

PRIOR APPLICATION NUMBER: 60/426,137
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NUMBER OF SEQ ID NOS: 673904
SOFTWARE: Microarray Probe Sequence Listing Generator V 1.1
SEQ ID NO 112898
LENGTH: 25
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SOPTWARE: Proprietary
SEQ ID NO 1230837
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Best Local Similarity 100.0%;
APPLICANT: Marshall, William
APPLICANT: Scaringe, Stephen
TITLE OF INVENTION: Punctional and Hyperfunctional sirna
FILE REFERENCE: 13499US
CURRENT APPLICATION UMBER: US/11/083,784
CURRENT FILING DATE: 2005-03-18
PRIOR FILING DATE: 2003-11-14
PRIOR FILING DATE: 2003-11-14
PRIOR PILING DATE: 2003-11-14
PRIOR FILING DATE: 2003-09-10
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APPLICANT: Khvobbva, Anastasia
APPLICANT: Reynolds, Angela
APPLICANT: Leake, Devin
APPLICANT: Marshall, William
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: RNA
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Similarity 61.5%;
8; Conservative
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Pred. No.
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Pred. No.
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Sequence 63320, Application US/10934048A

Publication No. US20060051770A1

GENERAL INFORMATION:
APPLICANT: Tanya Makeev
TITLE OF INVENTION: Methods of Genetic Analysis of Yeast
FILE REFERENCE: 3699
CURRENT APPLICATION NUMBER: US/10/934,048A
CURRENT FILING DATE: 2004-09-03
NUMBER OF SEQ ID NOS: 120855
SOFTWARE: Microarray Probe Sequence Listing Generator V 1.1
SEQ ID NO 68320
LENGTH: 25
TYPE: DNA
ORGANISM: S. pombe
US-10-934-048A-68320
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; TYPE: RNA
; ORGANISM: Homo mapiens
US-11-083-784-1230837
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US-10-934-048A-68320
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APPLICANT: Shiler, Kvuzat
PTITLE OF INVENTION: Bioinformatically detectable group of novel regulatory genes and
PTITLE OF INVENTION: uses thereof
FILE REFERENCE: 06087.0200.CPUS01
CURRENT APPLICATION NUMBER: US/10/310,914A
CURRENT FILING DATE: 2002-12-06
NUMBER OF SEQ ID NOS: 1388402
SOFTWARE: Patentin version 3.3
SEQ ID NO 224598
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 224598, Application US/10310914A Publication No. US20060003322A1 GENERAL INFORMATION:
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Best Local Similarity 61.5%;
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Best Local Similarity
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PRIOR FILING DATE: 2002-11-14
NUMBER OF SEQ ID NOS: 1591911
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TYPE: RNA
ORGANISM: Human
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                    CTCTTCTGGATAC 18
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                                                                            72.2%; Score 13; DB 9; 100.0%; Pred. No. 8e+02;
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Pred. No. 8e+02;
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Pred. No. 8e+02;
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RESULT 9
US-10-310-914A-439308/c
US-10-310-914A-439308, Application US/10310914A
; Publication No. US20060003322A1
; GENERAL INFORMATION:
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US-11-121-849-325988
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APPLICANT: Shiler, Kvuzat
TITLE OF INVENTION: Bioinformatically detectable group of novel regulatory genes and
TITLE OF INVENTION: uses thereof
FILE REFERENCE: 06087.0200.CPUS01
CURRENT APPLICATION NUMBER: US/10/310,914A
CURRENT FILING DATE: 2002-12-06
NUMBER OF SEQ ID NOS: 1388402
SOPTWARE: Patentin version 3.3
SEQ ID NO 224589
LENGTH: 27
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Best Local
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Best Local
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TITLE OF INVENTION: Methods of Genetic Analysis of Formalin Fixed Paraffin Embedded
TITLE OF INVENTION: Microarrays
FILE REFERENCE: 3684.1
CURRENT APPLICATION NUMBER: US/11/121,849
CURRENT FILING DATE: 2005-05-03
PRIOR APPLICATION NUMBER: 60/567,949
PRIOR APPLICATION NUMBER: 60/567,949
PRIOR FILING DATE: 2004-05-03
PRIOR FILING DATE: 2004-05-03
NUMBER OF SEQ ID NOS: 673904
APPLICANT: Bentwich, Isaac
APPLICANT: Shifer, Kvuzat
TITLE OP INVENTION: Bloinformatically detectable group of novel regulatory genes
TITLE OP INVENTION: uses thereof
FILE REPERENCE: 06087.0200.CPUS01
CURRENT APPLICATION NUMBER: US/10/310,914A
CURRENT FILING DATE: 2002-12-06
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9; Conser
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Pred. No. 8e+02;
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; Sequence 1226662, Application US/10310914A
; Publication No. US20060003322A1
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; Sequence 552172, Application US/10310914A; Publication No. US20060003322A1; GENERAL INFORMATION:
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US-10-310-914A-439308
                                                                                                                                                                                         NUMBER OF SEQ ID NOS: 1388402
SOFTWARE: PatentIn version 3.3
SEQ ID NO 1226662
                                                                              Query Match
Best Local
                                                               Matches
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Best Local
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SOFTWARE: PatentIn version 3.3
SEQ ID NO 439308
LENGTH: 18
                                                                                                                                                                                                                                    APPLICANT: Bentwich, Isaac APPLICANT: Shiler, Kruzat TITLE OF INVENTION: Bioinformatically detectable group of novel regulatory genes TITLE OF INVENTION: uses thereof FILE REFERENCE: 06087.0200.CPUS01 CURRENT APPLICATION NUMBER: US/10/310,914A CURRENT FILING DATE: 2002-12-06
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APPLICANT: Shiler, Kvuzat
APPLICANT: Shiler, Kvuzat
TITLE OF INVENTION: Bioinformatically detectable group of novel regulatory genes
TITLE OF INVENTION: uses thereof
FILE REFERENCE: 06087.0200.CPUS01
CURRENT FILING DATE: 2002-12-06
CURRENT FILING DATE: 2002-12-06
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Best Local Similarity
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                                                                                                                                            TYPE: RNA
ORGANISM: Human
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ORGANISM: Human
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                              CCCCTCTTCTGG 14
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                                                                             66.7%; Score 12; 100.0%; Pred. No.
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%; Pred. No. 2.8
0; Mismatches
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US-11-101-244-603439/c

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APPLICANT: Reynolds, Angela
APPLICANT: Leake, Devin
APPLICANT: Marshall, William
APPLICANT: Scaringe, Stephen
TITLE OF INVENTION: Functional and Hyperfunctional sirna
FILE REFERENCE: 13499US
CURRENT APPLICATION NUMBER: US/11/101,244
CURRENT FILING DATE: 2005-04-07
PRIOR APPLICATION NUMBER: 60/502,050
PRIOR APPLICATION NUMBER: 60/426,137
PRIOR FILING DATE: 2003-09-10
SOFTWARE: Proprietary
SOFTWARE: Proprietary
SEQ ID NO 815438
LENGTH: 19
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US-11-101-244-815438/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; TYPE: RNA; Homo sapiens US-11-101-244-603439
                                                                                                                                                                                                  ; ORGANISM: Homo, papiens
US-11-101-244-815438
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 603439, Application US/11101244 Publication No. US20050246794A1 GENERAL INFORMATION:
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LENGTH: 19
                                                                                                                         Query Match
Best Local &
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 815438, Application US/11101244 Publication No. US20050246794A1
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APPLICANT: Khvorova, Anastasia
APPLICANT: Reynolds, Angela
APPLICANT: Leake, Devin
APPLICANT: Marshall, William
APPLICANT: Scaringe, Stephen
TITLE OF INVENTION: Functional and Hyperfunctional sirNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Dharmacon, Inc.
APPLICANT: Khvorova, Anastasia
APPLICANT: Reynolds, Angela
APPLICANT: Leake, Devin
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CURRENT APPLICATION NUMBER: US/11/101,244
CURRENT FILING DATE: 2005-04-07
PRIOR APPLICATION NUMBER: 60/502,050
PRIOR FILING DATE: 2003-09-10
PRIOR APPLICATION NUMBER: 66/426,137
PRIOR APPLICATION NUMBER: 66/426,137
PRIOR PILING DATE: 2002-11-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQ ID NOS: 1591911
SOFTWARE: Proprietary
                                                                                                                    Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Hacult 66.7%; Score 12; DB 12; Local Similarity 100.0%; Pred. No. 2.8e+03; es 12; Conservative 0: Miamatair
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                                                    7 TCTTCTGGATAC 18
                                                                                               n 66.7%; Score 12; DB 12; Length 19;
Similarity 100.0%; Pred. No. 2.8e+03;
12; Conservative 0; Mismatches 0; Indels
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                                      S
                                                                                                                                                                          ; ORGANISM: Homo sapiens US-11-101-244-817168
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US-11-101-244-817168/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; ORGANISM: Homo sapiens
US-11-101-244-815537
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                                                                                                                                                                                                                                                                              CURRENT APPLICATION NUMBER: US/11/101,244
CURRENT FILING DATE: 2005-04-07
PRIOR APPLICATION NUMBER: 60/502,050
PRIOR FILING DATE: 2003-09-10
PRIOR APPLICATION NUMBER: 60/426,137
PRIOR APPLICATION NUMBER: 60/426,137
PRIOR FILING DATE: 2002-11-14
NUMBER OF SEQ ID NOS: 1591911
SOFTWARE: Proprietary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION NUMBER: 60/426,137
PRIOR FILING DATE: 2002-11-14
NUMBER OF SEQ ID NOS: 1591911
SOFTWARE: Proprietary
SEQ ID NO 815537
                                                                                                                                                                                                                   SEQ ID NO 817168
LENGTH: 19
TYPE: RNA
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APPLICANT: Khvorova, Anastasia
APPLICANT: Reynolds, Angela
APPLICANT: Reynolds, Angela
APPLICANT: Leake, Devin
APPLICANT: Marshall, William
APPLICANT: Scaringe, Stephen
TITLE OF INVENTION: Functional and Hyperfunctional sirNA
FILE REFERENCE: 13499US
CURRENT APPLICATION NUMBER: US/11/101,244
CURRENT FILING DATE: 2005-04-07
CURRENT FILING DATE: 2005-04-07
                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Marshall, William
APPLICANT: Scaringe, Stephen
TITLE OF INVENTION: Functional and Hyperfunctional sirNA
FILE REFERENCE: 13499US
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APPLICANT: Khvorova, Anastasia
APPLICANT: Reynolds, Angela
APPLICANT: Leake, Devin
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Local Similarity 100.0%; Pred. No.
les 12; Conservative 0; Mismatch
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                                                                                    ch 66.7%; Score 12; DB 12; l Similarity 100.0%; Pred. No. 2.8e+03; 12; Conservative 0; Mismatches 0;
                                  TCTTCTGGATAC 18
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b. 2.8e+03;
ches 0;
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Search completed: April 7, 2006, 18:19:23 Job time: 2275.35 secs --

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Perfect score:
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                derived by analysis of the total score distribution.
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1: /cgn2 { ptodata/1/pubpna/US07_PUBCOMB.seq: *

2: /cgn2 { ptodata/1/pubpna/US08_PUBCOMB.seq: *

3: /cgn2 { ptodata/1/pubpna/US09A_PUBCOMB.seq: *

4: /cgn2 { ptodata/1/pubpna/US09A_PUBCOMB.seq: *

4: /cgn2 { ptodata/1/pubpna/US09B_PUBCOMB.seq: *

5: /cgn2 { ptodata/1/pubpna/US10B_PUBCOMB.seq: *

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10: /cgn2 { ptodata/1/pubpna/US10B_PUBCOMB.seq: *
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         \begin{array}{c} \mathbf{7.7} \\ \mathbf{6.6} \\ \mathbf{6.6} \\ \mathbf{6.6} \\ \mathbf{6.7} \\ \mathbf{7.2} \\ \mathbf{2.2} \\ \mathbf{2.2
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Copyright (c) 1993 - 2006 Biocceleration Ltd.
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   7 US-10-659-948A-5

7 US-10-659-980A-5

7 US-10-719-956-117959

7 US-10-719-956-470164

7 US-10-719-956-511003

7 US-10-719-956-511802

7 US-10-719-956-511802

7 US-10-719-900-316050

7 US-10-719-900-574815

7 US-10-60-756-197597

8 US-10-60-756-197597

9 US-10-60-756-197597

9 US-10-60-756-197598

9 US-10-659-948A-6

9 US-10-659-948A-6

9 US-10-659-983A-6

9 US-10-659-983A-6
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Sequence 5, Appli
Sequence 5, Appli
Sequence 117959;
Sequence 193006,
Sequence 470164,
Sequence 511003,
Sequence 518629,
Sequence 611802,
Sequence 611802,
Sequence 197597,
Sequence 197597,
Sequence 197597,
Sequence 197598,
Sequence 6, Appli
Sequence 6, Appli
Sequence 6, Appli
Sequence 23, Appl
Sequence 243, Appl
Sequence 2513, Appl
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24 12 66.7 25 6 US-10-213-114-1 12-66.7 25 6 US-10-223-114-1 12-66.7 25 6 US-10-122-114-1 12-66.7 25 7 US-10-719-956-5244 12-7 12-7 12-7 12-7 12-7 12-7 12-7 12-7
66.7 25 7 US-10-66.7 25 8 US-1
6 US-10- 6 US-10- 7 US-10- 8 US-10- 8 US-10- 8 US-10- 8 US-10- 8 US-10-
US - 10 - 10 - 10 - 10 - 10 - 10 - 10 - 1

ALIGNMENTS

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CURRENT APPLICATION NUMBER: US/10/659,948A
CURRENT FILING DATE: 2003-09-10
PRIOR APPLICATION NUMBER: US 09/573,684
PRIOR FILING DATE: 2000-05-19
PRIOR PRILING DATE: 2002-09-19
PRIOR PILING DATE: 2002-09-19
PRIOR FILING DATE: 2002-09-19
PRIOR PILING DATE: 2002-09-19
PRIOR PILING DATE: 2002-09-19
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PRIOR PILING DATE: 2002-09-19
PRIOR FILING DATE: 2002-09-19
PRIOR FILING DATE: 2002-09-19
PRIOR FILING DATE: 2002-09-19
NUMBER OF SEQ ID NOS: 23
SOPTWARE: PatentIn version 3.2
SEQ ID NO 5
LENGTH: 18
CREATURE: Artificial
RESULT 2
US-10-659-980A-5
; Sequence 5, Application US/10659980A
; Publication No. US20040106133A1
; GENERAL INFORMATION:
; APPLICANT: Hovanec, Timothy A
; TITLE OF INVENTION: Method for Detecting Ammonia-Oxidizing Bacteria
; FILE REFERENCE: 81289-2284781
; FILE REFERENCE: 81289-2181
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; OTHER INFORMATION: Oligonucleotide Probe
US-10-659-948A-5
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US-10-659-948A-5
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Best Local
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                                                                                                                                                                                                                                                                         1 CCCCCCTCTTCTGGATAC 18
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100.0%; Pred. No. 2
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Sequence 5, Application US/10659983A
Publication No. US20040157313A1
GENERAL INFORMATION:
APPLICANT: Hovanec, Timothy A
TITLE OF INVENTION: Ammonia-Oxidizing Bacteria
FILE REFERENCE: 8129-284779
CURRENT APPLICATION UNMER: US/10/659,983A
CURRENT FILING DATE: 2003-09-10
PRIOR APPLICATION NUMBER: US 99/573,684
PRIOR FILING DATE: 2000-05-19
PRIOR FILING DATE: 2000-05-19
PRIOR FILING DATE: 2002-09-19
PRIOR APPLICATION NUMBER: US 60/386,218
PRIOR APPLICATION NUMBER: US 60/386,218
PRIOR APPLICATION NUMBER: US 60/386,219
                                                              RESULT 4
US-10-719-956-117959/c
; Sequence 117959, Application US/10719956
; Publication No. US20040146910A1
; GENERAL INFORMATION:
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US-10~659-983A-5
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                                                                                                                                                                                                                                                                                                                                                                                                    ; OTHER INFORMATION: Oligonucleotide Probe US-10-659-983A-5
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Best Local :
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SEQ ID NO 5
LENGTH: 18
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Best Local Similarity
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PRIOR FILING DATE: 2000-05-19
PRIOR APPLICATION NUMBER: US 60/386,217
PRIOR FILING DATE: 2002-09-19
PRIOR APPLICATION NUMBER: US 60/386,218
PRIOR FILING DATE: 2002-09-19
PRIOR FILING DATE: 2002-09-19
                    APPLICANT: Xue Mei Zhou
TITLE OF INVENTION: Methods of Genetic Analysis of Rat
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SOFTWARE: PatentIn version 3.2
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PRIOR FILING DATE: 2002-09-19
NUMBER OF SEQ ID NOS: 23
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                                                                                                                                                                                                                                                                                                                                                                                                                                                ORGANISM: Artificial FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: DNA
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ORGANISM: Artificial
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REFERENCE: 3527.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         18;
                                                                                                                                                                                                                                                                 1 CCCCCCTCTTCTGGATAC 18
                                                                                                                                                                                                                                                                                                                                  Similarity
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Pred. No. 2.2;
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Sequence 470164, Application US/10719956

Publication No. US20040146910A1

GENERAL INFORMATION:
APPLICANT: Xue Mei Zhou
TITLE OF INVENTION: Methods of Genetic Analysis of Rat
FILE REFERENCE: 3527.1

CURRENT APPLICATION NUMBER: US/10/719,956

CURRENT FILING DATE: 2003-11-20
PRIOR APPLICATION NUMBER: 60/427,836
PRIOR APPLICATION NUMBER: 60/427,836
PRIOR FILING DATE: 2002-11 20
NUMBER OF SEQ ID NOS: 699466

SOFTWARE: Microarray Probe Sequence Listing Generator V 1.1
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US-10-719-956-193006
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                                                                     ; ORGANISM: Rattus norvegicus US-10-719-956-470164
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US-10-719-956-470164
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US-10-719-956-193006
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                                                                                                          SEQ ID NO 470164
LENGTH: 25
TYPE: DNA
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PRIOR FILING DATE: 2002 11 20
NUMBER OF SEQ ID NOS: 699466
SOFTWARE: Microarray Probe Sequence Listing Generator V 1.1
SEQ ID NO 117959
Query Match 72.2*; So
Best Local Similarity 100.0%; P
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Publication No. US20040146910A1
GENERAL INFORMATION:
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LENGTH: 25
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CURRENT FILING DATE: 2003-11-20

PRIOR APPLICATION NUMBER: 60/427,836

PRIOR FILING DATE: 2002 11 20

NUMBER OF SEQ ID NOS: 699466

SOFTWARE: Microarray Probe Sequence Listing Generator V 1.1
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TITLE OF INVENTION: Methods of Genetic Analysis
FILE REFERENCE: 3527.1
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CURRENT FILING DATE: 2003-11-20
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TYPE: DNA
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100.0%; Pr
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   Score 13; DB 7; L
Pred. No. 1.3e+03;
0; Mismatches 0;
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Pred. No. 1.3e+03;
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US-10-719-956-558629
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US-10-719-956-558629
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US-10-719-956-511003
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US-10-719-956-511003
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Sequence 611802, Application US/10719956
Publication No. US20040146910A1
GENERAL INFORMATION:
APPLICANT: Xue Mei Zhou
TITLE OF INVENTION: Methods of Genetic Analysis of Rat
FILE REFERENCE: 3527.1
CURRENT APPLICATION NUMBER: US/10/719,956
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CURRENT FILING DATE: 2003-11-20
PRIOR APPLICATION NUMBER: 60/427,836
PRIOR FILING DATE: 2002 11 20
NUMBER OF SEQ ID NOS: 699466
SOFTWARE: Microarray Probe Sequence Listing Generator V 1.1
SEQ ID NO 511003
                                                                                                                                                                                                                                                                                               Matches
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Best Local :
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Best Local Similarity
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Publication No. US20040148910A1
GENERAL INFORMATION:
APPLICANT: Xue Mei Zhou
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APPLICANT: Xue Med Zhou

TITLE OF INVENTION: Methods of Genetic Analysis of Rat

FILE REFERENCE: 3527.1

CURRENT APPLICATION NUMBER: US/10/719,956

CURRENT FILING DATE: 2003-11-20

PRIOR APPLICATION NUMBER: 60/427,836

PRIOR APPLICATION NUMBER: 2002 11 20

RIOR PILING DATE: 2002 11 20

RUMBER OF SEQ ID NOS: 699466

SOFTWARE: Microarray Probe Sequence Listing Generator V 1.1
                                                                                                                                                                                                                                                                                                                                                                                                             LENGTH: 25
TYPE: DNA
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TYPE: DNA
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100.0%; Pred. No. 1.3e+03;
7ative 0; Mismatches 0;
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100.0%; Pred. No. 1.3e+03;
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US-10-719-900-574815
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US-10-719-900-574815
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; TYPE: DNA
; ORGANIEM: Rattus norvegicus
US-10-719-956-611802
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                                                                                                                                APPLICANT: Xue Mei Zhou

APPLICANT: Xue Mei Zhou

TITLE OF INVENTION: Methods of Genetic Analysis of Mouse

FILE REFERENCE: 3528.1

CURRENT APPLICATION NUMBER: US/10/719,900

CURRENT FILING DATE: 2003-11-20

PRIOR FILING DATE: 2003-11-20

PRIOR FILING DATE: 2002-11-20

NUMBER OF SEQ ID NOS: 982914

SOFTWARE: Microarray Probe Sequence Listing Generator V 1.1

SEQ ID NO 574815
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Xue Mei Zhou

TITLS OF INVENTION: Methods of Genetic Analysis of Mouse
FILE REFERENCE: 5528.1

CURRENT APPLICATION NUMBER: US/10/719,900

CURRENT FILING DATE: 2003-11-20
PRIOR APPLICATION NUMBER: 60/427,808
PRIOR FILING DATE: 2002 11 20

NUMBER OF SEQ ID NOS: 982914

SOFTWARE: Microarray Probe Sequence Listing Generator V 1.1

SEQ ID NO 316050

LENGTH: 25
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Best Local Similarity
Matches 13; Conserval
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Publication No. US20050026164A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
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PRIOR FILING DATE: 2002 11 20
NUMBER OF SEQ ID NOS: 699466
SOPTWARE: Microarray Probe Sequence Listing Generator V 1.1
SEQ ID NO 611802
                                        Query Match
   Matches
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                                                                                         LENGTH: 25
TYPE: DNA
ORGANISM: Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: DNA
ORGANISM: Mus musculus
   Local Similarity hes 13; Conserv
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   Conservative
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72.2%; Score 13; DB 8; L
100.0%; Pred. No. 1.3e+03;
tive 0; Mismatches 0;
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Sequence 197597, Application US/11060756

Publication No. US20050221354A1

GENERAL INFORMATION:

APPLICANT: Wyeth
APPLICANT: Wounts, William Martin
APPLICANT: Wounts, William Martin
TITLE OF INVENTION: Target Genes
FILE OF INVENTION TARGET Genes
FILE REFERENCE: AM101083 (031896-042000)
CURRENT APPLICATION NUMBER: US/11/060,756
CURRENT APPLICATION NUMBER: US/11/060,756
UNMBER OF SEQ ID NOS: 303284
SOFTWARE: Patentin version 3.2
SEQ ID NO 197597
LENGTH: 25
TYPE: DNA
TYPE: D
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Mounts, William Martin
TITLE OF INVENTION: Nucleic Acid Arrays for Mor
TITLE OF INVENTION: Target Genes
FILE REFERENCE: AM101083 (031896-042000)
CURRENT APPLICATION NUMBER: US/11/060,756
CURRENT FILING DATE: 2005-02-18
NUMBER OF SEQ ID NOS: 303284
SOFTMARE: Patentin version 3.2
SEQ ID NO 197598
LENGTH: 25
TYPE: DNA
ORGANISM: probe
US-11-060-756-197598
                                                                                                          US-10-459-876-17
Sequence 17, Application US/10459876
Publication No. US20030208060A1
GENERAL INFORMATION:
APPLICANT: Gianturco, S.H.
APPLICANT: Bradley, W.A.
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Best Local Similarity
Watches 13; Conserve
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US-11-060-756-197597/c
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Best Local (
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APPLICANT: Mounts
APPLICANT: Bradley, W.A.
TITLE OF INVENTION: DAMA Encoding Human apoB48R: A Monocyte-Macrophage
TITLE OF INVENTION: Apolipoprotein B48 Receptor Gene and Protein
FILE REFERENCE: D5880CIP
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Search completed: April Job time: 350.784 Becs

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2006, 09:17:27

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CURRENT APPLICATION NUMBER: US/10/459,876
CURRENT FILING DATE: 2003-06-12
PRIOR APPLICATION NUMBER: US/09/583,610
PRIOR FILING DATE: 2000-05-31
PRIOR APPLICATION NUMBER: US 09/130,242
PRIOR FILING DATE: 1998-08-06
NUMBER OF SEQ ID NOS: 22
SEQ ID NO 17
LENGTH: 17
TYPE: DNA
ORGANISM: Artificial Sequence
PEATURE:
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US-10-659-948A-6/c
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                             á
                                                                                                                                                ; OTHER INFORMATION: PCR Primer US-10-659-948A-6
                                                                                                                                                                            SEQ ID NO 6
LENGTH: 18
TYPE: DNA
ORGANISM: Artificial
PROGRATURE:
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CURRENT APPLICATION NUMBER: US/10/659,948A

CURRENT FILING DATE: 2003-09-10

PRIOR APPLICATION NUMBER: US 09/573,684

PRIOR FILING DATE: 2000-05-19

PRIOR APPLICATION NUMBER: US 60/386,217

PRIOR FILING DATE: 2002-09-19

PRIOR PRIOR PRICATION NUMBER: US 60/386,218

PRIOR PRIOR PRICATION NUMBER: US 60/386,218

PRIOR FILING DATE: 2002-09-19

PRIOR APPLICATION NUMBER: US 60/386,219

PRIOR PILING DATE: 2002-09-19

NUMBER OF SEQ ID NOS: 23

SOFTWARE: Patentin version 3.2
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                                                                     Query Match
Best Local Similarity
Matches 12; Conserv
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Publication No. US20040101946A1
GENERAL INFORMATION:
APPLICANT: Hovanec, Timothy A
TITLE OF INVENTION: Method of Using Ammonia-Oxidizing Bacteria
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Pred. No.
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